Fig. 1

	A	B	C	D	E	F	G	H
1	211296_x_at	ubiquitin C	Hs.183704.0	gb:AB009010.1	UbC1		3.70	0.0199
2	200007_at	signal recognition particle 14kD (homologous Alu RNA-binding protein)	Hs.180394.0	gb:NM_003134.1	SRP14	15	3.02	0.0236
3	211943_x_at	tumor protein, translationally-controlled 1	Hs.279860.3	gb:AL565449_RC		12	3.58	0.0263
4	200633_at	ubiquitin B	Hs.183842.0	gb:NM_018955.1	UBB	17	3.62	0.0263
5	201429_s_at	ribosomal protein L37a	Hs.5566.0	gb:NM_000998.1	RPL37A	2	3.80	0.0264
6	200630_x_at	SET translocation (myeloid leukemia-associated)	Hs.145279.0	gb:AV702810		9	2.82	0.0270
7	212661_x_at	peptidylprolyl isomerase A (cyclophilin A)	Hs.182937.2	gb:BE731738		21	3.51	0.0274
8	201273_s_at	signal recognition particle 9kD	Hs.75975.0	gb:NM_003133.1	SRP9	1	2.89	0.0281
9	200870_at	unr-interacting protein	Hs.3727.0	gb:NM_007178.1	UNRIP	12	2.78	0.0281
10	200097_s_at	heterogeneous nuclear ribonucleoprotein K	Hs.129548.1	gb:AI701949_RC		9	2.73	0.0286
11	211956_s_at	putative translation initiation factor	Hs.150580.3	gb:BF246436		2	3.45	0.0290
12	214288_s_at	proteasome (prosome, macropain) subunit, beta type, 1	Hs.75748.1	gb:W86293_RC		6	2.93	0.0295
13	201293_x_at	peptidylprolyl isomerase A (cyclophilin A)	Hs.182937.0	gb:NM_021130.1	PPIA	7	3.50	0.0301
14	208996_s_at	polymerase (RNA) II (DNA directed) polypeptide C (33kD)	Hs.79402.0	gb:BC000409.1		16	2.39	0.0302
15	207320_x_at	staufen (Drosophila, RNA-binding protein)	Hs.6113.1	gb:NM_004602.1	STAU	20	2.75	0.0302
16	208780_x_at	VAMP (vesicle-associated membrane protein)-associated protein A (33kD)	Hs.9006.0	gb:AF154847.1	VAMP	18	2.97	0.0310
17	212284_x_at	hypothetical protein FLJ20030	Hs.326456.1	gb:BG498776		13	3.76	0.0312
18	201257_x_at	ribosomal protein S3A	Hs.77039.0	gb:NM_001006.1	RPS3A	4	3.69	0.0315
19	211628_x_at			gb:J04755.1	FTHP1	6	3.51	0.0316
20	212643_at	Homo sapiens cDNA: FLJ21927 fis, clone HEP04178, highly similar to HS	Hs.81360.1	gb:AI671747_RC		14	2.76	0.0317
21	211978_x_at	KIAA1228 protein	Hs.267690.2	gb:AI708767_RC		7	3.52	0.0318
22	212869_x_at	hypothetical protein FLJ20030	Hs.326456.2	gb:AI721229_RC		12	3.83	0.0318
23	212391_x_at	ribosomal protein S3A	Hs.77039.1	gb:AI925635_RC		15	3.67	0.0319
24	213037_x_at	staufen (Drosophila, RNA-binding protein)	Hs.6113.5	gb:AJ132258.1	staufen	20	2.72	0.0320
25	49878_at			gb:AA523441			2.47	0.0321
26	200021_at	cofilin 1 (non-muscle)	Hs.180370.0	gb:NM_005507.1	CFL1	11	3.56	0.0328
27	202028_s_at	ribosomal protein L38	Hs.2017.0	gb:BC000603.1_RC		17	2.49	0.0328
28	213738_s_at	ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, i	Hs.155101.3	gb:AI587323_RC		2	3.19	0.0328
29	213084_x_at	cadherin 1, type 1, E-cadherin (epithelial)	Hs.194657.4	gb:BF125158		3	3.70	0.0329
30	201225_s_at	SerArg-related nuclear matrix protein (plenty of prolines 101-like)	Hs.18192.0	gb:NM_005839.1	SRM160	1	2.48	0.0334
31	208781_x_at	sorting nexin 3	Hs.12102.0	gb:AF062483.1		6	2.87	0.0339

Fig. 2

	A	B	C	D	E	F	G	H
32	200718_s_at	transcription elongation factor B (SIII), polypeptide 1-like	Hs.171626.0	gb:AA927664_RC		5	3.25	0.0339
33	211378_x_at	KIAA1228 protein	Hs.267690.0	gb:BC001224.1			3.49	0.0339
34	208834_x_at	cadherin 1, type 1, E-cadherin (epithelial)	Hs.194657.1	gb:BC001865.1		17	3.72	0.0340
35	213881_x_at	SMT3 (suppressor of mif two 3, yeast) homolog 2	Hs.180139.1	gb:AI971724_RC		17	3.06	0.0340
36	200627_at	unactive progesterone receptor, 23 kD	Hs.278270.0	gb:BC003005.1		12	2.86	0.0341
37	200077_s_at			gb:D87914.1	hAZ-brain	19	3.28	0.0343
38	200030_s_at	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member	Hs.78713.0	gb:NM_002635.1	SLC25A3	12	3.22	0.0344
39	208842_s_at	DKFZP434D156 protein	Hs.6880.1	gb:W93787_RC		2	2.56	0.0344
40	206559_x_at	eukaryotic translation elongation factor 1 alpha 1-like 14	Hs.274466.0	gb:NM_001403.1	EEF1A1L	6	3.71	0.0345
41	202300_at	hepatitis B virus x-interacting protein (9.6kD)	Hs.80464.0	gb:NM_006402.1	XIP	1	2.66	0.0347
42	207243_s_at	calmodulin 2 (phosphorylase kinase, delta)	Hs.182278.0	gb:NM_001743.1	CALM2	12	3.46	0.0347
43	221741_s_at	hypothetical protein FLJ20391	Hs.11747.1	gb:AL096828		20	2.42	0.0347
44	201553_s_at	lysosomal-associated membrane protein 1	Hs.150101.0	gb:NM_005561.2	LAMP1	13	3.22	0.0347
45	208826_x_at	histidine triad nucleotide-binding protein	Hs.256697.1	gb:U27143.1		5	3.07	0.0348
46	214737_x_at	heterogeneous nuclear ribonucleoprotein C (C1C2)	Hs.182447.3	gb:AV725195		14	2.76	0.0350
47	210453_x_at	ATP synthase, H+ transporting, mitochondrial F1F0, subunit g	Hs.107476.2	gb:AL050277.1	DKFZp56	11	3.05	0.0351
48	221791_s_at	hypothetical protein	Hs.171774.1	gb:BG167522		3	3.12	0.0351
49	221476_s_at	ribosomal protein L15	Hs.74267.0	gb:AF279903.1	EC45	3	3.22	0.0353
50	200963_x_at	ribosomal protein L31	Hs.184014.0	gb:NM_000993.1	RPL31	2	3.55	0.0353
51	65635_at			gb:/AL044097			2.40	0.0355
52	200717_x_at	ribosomal protein L7	Hs.153.0	gb:NM_000971.1	RPL7	8	3.57	0.0355
53	200059_s_at	ras homolog gene family, member A	Hs.77273.0	gb:BC001360.1		3	3.16	0.0356
54	217720_at	16.7Kd protein	Hs.180859.0	gb:NM_016139.1	LOC5114	7	3.19	0.0357
55	208948_s_at	staufen (Drosophila, RNA-binding protein)	Hs.6113.4	gb:BC000830.1		20	2.64	0.0358
56	200912_s_at	eukaryotic translation initiation factor 4A, isoform 2	Hs.173912.0	gb:NM_001967.2	EIF4A2	3	3.34	0.0359
57	200893_at	splicing factor, arginineserine-rich (transformer 2 Drosophila homolog) 10	Hs.30035.0	gb:NM_004593.1	SFRS10	3	2.71	0.0361
58	202029_x_at	ribosomal protein L38	Hs.2017.0	gb:NM_000999.1	RPL38	17	3.46	0.0361
59	208825_x_at	ribosomal protein L23a	Hs.184776.1	gb:U43701.1		17	3.72	0.0362
60	210648_x_at	sorting nexin 3	Hs.12102.1	gb:AB047360.1	SNX 3A	6	3.15	0.0363
61	217830_s_at	p47	Hs.12865.0	gb:AL109658		20	2.28	0.0364
62	208724_s_at	RAB1, member RAS oncogene family	Hs.3642.2	gb:BC000905.1		2	3.03	0.0365

Fig. 3

	A	B	C	D	E	F	G	H
63	211765_x_at			gb:BC005982.1		7	3.48	0.0365
64	212261_at	trinucleotide repeat containing 15	Hs.323317.0	gb:AB014542.1	KIAA064	2	2.26	0.0365
65	221494_x_at	muscle specific gene	Hs.283781.0	gb:AF085358.1		19	2.86	0.0368
66	200018_at	ribosomal protein S13	Hs.165590.0	gb:NM_001017.1	RPS13	11	3.45	0.0368
67	217732_s_at	integral membrane protein 2B	Hs.239625.0	gb:AF092128.1		13	3.25	0.0370
68	201754_at	cytochrome c oxidase subunit VIc	Hs.74649.0	gb:NM_004374.1	COX6C	18	3.10	0.0371
69	217801_at	ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit	Hs.177530.0	gb:NM_006886.1	ATP5E	20	2.94	0.0371
70	200039_s_at	proteasome (prosome, macropain) subunit, beta type, 2	Hs.1390.0	gb:NM_002794.1	PSMB2	1	2.61	0.0372
71	200773_x_at	prothymosin, alpha (gene sequence 28)	Hs.250655.0	gb:NM_002823.1	PTMA	5	3.07	0.0372
72	213179_at	ESTs	Hs.293687.0	gb:BG289914		2	2.13	0.0373
73	200819_s_at	ribosomal protein S15	Hs.133230.0	gb:NM_001018.1	RPS15	19	3.52	0.0373
74	200006_at	RNA-binding protein regulatory subunit	Hs.10958.0	gb:NM_007262.1	DJ-1	1	3.10	0.0374
75	214789_x_at	Splicing factor, arginineserine-rich, 46kD	Hs.155160.2	gb:AA524274_RC		11	2.56	0.0374
76	200019_s_at	Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expr	Hs.177415.0	gb:NM_001997.1	FAU	11	3.33	0.0376
77	216520_s_at	tumor protein, translationally-controlled 1	Hs.279860.5	gb:AF072098			3.66	0.0376
78	212130_x_at	putative translation initiation factor	Hs.150580.4	gb:AL537707		X	3.32	0.0377
79	208904_s_at	ribosomal protein S28	Hs.153177.1	gb:BC000354.1		19	3.50	0.0377
80	200084_at	small acidic protein	Hs.78050.1	gb:BE748698_RC		14	2.65	0.0378
81	208738_x_at	SMT3 (suppressor of mif two 3, yeast) homolog 2	Hs.180139.0	gb:AK024823.1		17	2.76	0.0378
82	221607_x_at	actin, gamma 1	Hs.14376.1	gb:BC001920.1		1	3.51	0.0378
83	216221_s_at	pumilio (Drosophila) homolog 2	Hs.6151.1	gb:D87078.2	KIAA023	2	2.57	0.0379
84	200080_s_at	H3 histone, family 3A	Hs.181307.5	gb:AI955655_RC		2	3.14	0.0381
85	207573_x_at	ATP synthase, H+ transporting, mitochondrial F1F0, subunit g	Hs.107476.0	gb:NM_006476.1	ATP5JG	11	2.95	0.0381
86	200016_x_at	heterogeneous nuclear ribonucleoprotein A1	Hs.249495.0	gb:NM_002136.1	HNRPA1	18	3.28	0.0382
87	212227_x_at	putative translation initiation factor	Hs.150580.2	gb:W67644_RC		17	3.30	0.0384
88	212988_x_at	actin, gamma 1	Hs.14376.3	gb:AL515810_RC		3	3.52	0.0385
89	202698_x_at	cytochrome c oxidase subunit IV	Hs.113205.0	gb:NM_001861.1	COX4	16	3.25	0.0385
90	200863_s_at	RAB11A, member RAS oncogene family	Hs.75618.0	gb:AI215102_RC		15	2.77	0.0386
91	200012_x_at	ribosomal protein L21	Hs.184108.0	gb:NM_000982.1	RPL21	10	3.38	0.0387
92	200004_at	eukaryotic translation initiation factor 4 gamma, 2	Hs.183684.0	gb:NM_001418.1	EIF4G2	11	3.07	0.0388
93	200645_at	GABA(A) receptor-associated protein	Hs.7719.0	gb:NM_007278.1	GABARA	17	3.14	0.0389

Fig. 4

	A	B	C	D	E	F	G	H
94	217491_x_at	cytochrome c oxidase subunit VIIc pseudogene 1	Hs.278494.0	gb:AF042165			3.01	0.0389
95	201240_s_at	KIAA0102 gene product	Hs.77665.0	gb:NM_014752.1	KIAA010	11	2.84	0.0392
96	208718_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 17 (72kD)	Hs.6179.1	gb:Z97056		22	3.19	0.0392
97	208746_x_at	ATP synthase, H+ transporting, mitochondrial F1F0, subunit g	Hs.107476.1	gb:AF070655.1		11	3.01	0.0393
98	217982_s_at	MORF-related gene 15	Hs.6353.0	gb:NM_006791.1	MRG15	1	3.15	0.0393
99	200740_s_at	SMT3 (suppressor of mif two 3, yeast) homolog 1	Hs.85119.0	gb:NM_006936.1	SMT3H1	21	2.77	0.0394
100	212581_x_at	glyceraldehyde-3-phosphate dehydrogenase	Hs.169476.1	gb:BE561479		12	3.64	0.0396
101	207132_x_at	prefoldin 5	Hs.288856.0	gb:NM_002624.1	PFDN5	12	3.00	0.0397
102	201550_x_at	actin, gamma 1	Hs.14376.0	gb:NM_001614.2	ACTG1	1	3.52	0.0397
103	212857_x_at	activated RNA polymerase II transcription cofactor 4	Hs.74861.1	gb:BG231551_RC		16	2.81	0.0400
104	213477_x_at	PRO2047 protein	Hs.284136.2	gb:AL515273_RC		9	3.63	0.0400
105	200068_s_at	calnexin	Hs.155560.0	gb:M94859.1		UL	3.06	0.0402
106	200741_s_at	ribosomal protein S27 (metallopanstimulin 1)	Hs.195453.0	gb:NM_001030.1	RPS27	1	3.57	0.0403
107	211961_s_at	hypothetical protein PRO2706	Hs.237955.3	gb:AK000826.1		3	2.64	0.0405
108	221495_s_at	KIAA1049 protein	Hs.227835.0	gb:AF322111.1		NA	2.66	0.0405
109	218048_at	BUP protein	Hs.35660.0	gb:NM_012071.1	BUP	10	2.31	0.0405
110	211970_x_at	actin, gamma 1	Hs.14376.4	gb:BG026805		X	3.51	0.0406
111	202244_at	proteasome (prosome, macropain) subunit, beta type, 4	Hs.89545.0	gb:NM_002796.1	PSMB4	1	2.72	0.0406
112	200594_x_at	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	Hs.103804.0	gb:NM_004501.1	HNRPU	1	2.75	0.0406
113	211995_x_at	actin, gamma 1	Hs.14376.9	gb:AL567820_RC		1	3.53	0.0406
114	210908_s_at	prefoldin 5	Hs.288856.2	gb:AB055804.1	mm-1	12	2.89	0.0406
115	211983_x_at	actin, gamma 1	Hs.14376.2	gb:BE741683		3	3.53	0.0408
116	200065_s_at	ADP-ribosylation factor 1	Hs.74571.0	gb:AF052179.1		1	3.01	0.0411
117	201492_s_at	ribosomal protein L41	Hs.324406.0	gb:NM_021104.1	RPL41	12	3.68	0.0412
118	200780_x_at	guanine nucleotide binding protein (G protein), alpha stimulating activity p	Hs.273385.0	gb:NM_000516.2	GNAS1	20	3.49	0.0412
119	221943_x_at	ribosomal protein L38	Hs.2017.1	gb:AW303136_RC		17	2.68	0.0412
120	201817_at	KIAA0010 gene product	Hs.155287.0	gb:NM_014671.1	KIAA001	7	2.14	0.0413
121	200603_at	protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific e	Hs.183037.0	gb:AL050038.1		17	2.83	0.0413
122	200061_s_at	ribosomal protein S24	Hs.180450.1	gb:BC000523.1		10	3.45	0.0413
123	200067_x_at	sorting nexin 3	Hs.12102.0	gb:AL078596		6	2.86	0.0414
124	31826_at						2.24	0.0414

Fig. 5

	A	B	C	D	E	F	G	H
125	200981_x_at	neuroendocrine secretory protein 55	Hs.113368.0	gb:NM_016592.1	NESP55	20	3.49	0.0414
126	212266_s_at	splicing factor, arginineserine-rich 5	Hs.166975.3	gb:AW084582_RC		14	2.77	0.0416
127	212042_x_at	ribosomal protein L7	Hs.153.1	gb:BG389744		1	3.27	0.0418
128	202021_x_at	putative translation initiation factor	Hs.150580.0	gb:AF083441.1		17	3.30	0.0418
129	203752_s_at	jun D proto-oncogene	Hs.2780.0	gb:NM_005354.2	JUND	19	3.06	0.0419
130	207974_s_at	S-phase kinase-associated protein 1A (p19A)	Hs.227950.0	gb:NM_006930.1	SKP1A		2.84	0.0420
131	200926_at	ribosomal protein S23	Hs.3463.0	gb:NM_001025.1	RPS23	5	3.55	0.0420
132	203406_at	microfibrillar-associated protein 1	Hs.61418.0	gb:NM_005926.1	MFAP1	15	2.32	0.0420
133	213583_x_at	tRNA isopentenylpyrophosphate transferase	Hs.288036.2	gb:BE964125_RC		13	3.49	0.0420
134	218467_at	x 003 protein	Hs.3726.0	gb:NM_020232.1	MDS003	18	2.65	0.0421
135	214003_x_at	ribosomal protein S20	Hs.8102.1	gb:BF184532		2	3.43	0.0421
136	217225_x_at	pM5 protein	Hs.227823.1	gb:AL512687.1	DKFZp54	16	2.98	0.0421
137	213047_x_at	SET translocation (myeloid leukemia-associated)	Hs.145279.2	gb:AI278616		9	2.72	0.0421
138	211542_x_at	ribosomal protein S10	Hs.76230.1	gb:BC004334.1			3.69	0.0422
139	201134_x_at	cytochrome c oxidase subunit VIIc	Hs.3462.0	gb:NM_001867.1	COX7C	5	3.07	0.0422
140	210646_x_at	ribosomal protein L13a	Hs.119122.1	gb:BC001675.1		10	3.72	0.0423
141	204102_s_at	eukaryotic translation elongation factor 2	Hs.75309.0	gb:NM_001961.1	EEF2	19	3.39	0.0423
142	212971_at	cysteinyl-tRNA synthetase	Hs.159604.1	gb:AI769685_RC		11	2.66	0.0423
143	200631_s_at	SET translocation (myeloid leukemia-associated)	Hs.145279.0	gb:NM_003011.1	SET	9	2.70	0.0424
144	212273_x_at	neuroendocrine secretory protein 55	Hs.113368.2	gb:AI591100_RC		20	3.45	0.0424
145	201500_s_at	protein phosphatase 1, regulatory (inhibitor) subunit 11	Hs.82887.0	gb:NM_021959.1	PPP1R11	6	2.66	0.0424
146	201812_s_at	6.2 kd protein	Hs.112318.0	gb:NM_019059.1	LOC5454	7	3.13	0.0424
147	201166_s_at	pumilio (Drosophila) homolog 1	Hs.153834.0	gb:NM_014676.1	PUM1	1	2.66	0.0425
148	201997_s_at	KIAA0929 protein Msx2 interacting nuclear target (MINT) homolog	Hs.184245.0	gb:NM_015001.1	KIAA092	1	2.42	0.0425
149	206323_x_at	oligophrenin 1	Hs.128824.0	gb:NM_002547.1	OPHN1	X	2.62	0.0425
150	208640_at	ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding p	Hs.173737.0	gb:BG292367		7	3.18	0.0427
151	200786_at	proteasome (prosome, macropain) subunit, beta type, 7	Hs.118065.0	gb:NM_002799.1	PSMB7	9	2.60	0.0427
152	203667_at	tubulin-specific chaperone a	Hs.24930.0	gb:NM_004607.1	TBCA	5	2.73	0.0427
153	201778_s_at	KIAA0494 gene product	Hs.62515.0	gb:NM_014774.1	KIAA049	1	2.70	0.0428
154	204892_x_at	eukaryotic translation elongation factor 1 alpha 1	Hs.181165.0	gb:NM_001402.1	EEF1A1	6	3.69	0.0428
155	201575_at	SKI-INTERACTING PROTEIN	Hs.79008.0	gb:NM_012245.1	SNW1	14	2.35	0.0429

Fig. 6

	A	B	C	D	E	F	G	H
156	211069_s_at			gb:BC006462.1		2	2.48	0.0429
157	211755_s_at			gb:BC005960.1		1	3.02	0.0429
158	200093_s_at	histidine triad nucleotide-binding protein	Hs.256697.2	gb:N32864_RC		5	2.77	0.0429
159	219547_at	COX15 (yeast) homolog, cytochrome c oxidase assembly protein	Hs.226581.0	gb:NM_004376.1	COX15	10	2.09	0.0429
160	217927_at	HSPC033 protein	Hs.11125.0	gb:NM_014041.1	HSPC033	3	2.91	0.0430
161	217898_at	chromosome 11 hypothetical protein ORF3	Hs.4245.0	gb:NM_020154.1	LOC5685	15	2.43	0.0430
162	212043_at	Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	Hs.110771.0	gb:W72053_RC		2	2.69	0.0430
163	208742_s_at	sin3-associated polypeptide, 18kD	Hs.23964.0	gb:U78303.1		13	2.67	0.0431
164	200041_s_at	HLA-B associated transcript-1	Hs.55296.0	gb:NM_004640.1	D6S81E	6	2.82	0.0431
165	207721_x_at	histidine triad nucleotide-binding protein	Hs.256697.0	gb:NM_005340.1	HINT	5	3.03	0.0432
166	49327_at			gb:AI492888			2.49	0.0432
167	213214_x_at	actin, gamma 1	Hs.14376.5	gb:AW190090_RC		2	3.50	0.0432
168	203034_s_at	ribosomal protein L27a	Hs.76064.0	gb:NM_000990.1	RPL27A	6	3.58	0.0432
169	213614_x_at	PRO2047 protein	Hs.284136.5	gb:BE786672		7	3.58	0.0434
170	211999_at	H3 histone, family 3B (H3.3B)	Hs.180877.1	gb:Z48950		17	3.06	0.0435
171	200032_s_at	ribosomal protein L9	Hs.157850.0	gb:NM_000661.1	RPL9	4	3.50	0.0436
172	202117_at	Rho GTPase activating protein 1	Hs.138860.0	gb:BG468434			2.70	0.0436
173	212788_x_at	ferritin, light polypeptide	Hs.111334.1	gb:BG537190		20	3.67	0.0437
174	213453_x_at	glyceraldehyde-3-phosphate dehydrogenase	Hs.169476.4	gb:BF689355_RC		13	3.48	0.0439
175	213867_x_at	actin, beta	Hs.288061.2	gb:AA809056_RC		6	3.64	0.0440
176	210996_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.79474.1	gb:U43430.1		17	2.78	0.0441
177	208845_at	voltage-dependent anion channel 3	Hs.7381.0	gb:BC002456.1		8	2.89	0.0441
178	217740_x_at	ribosomal protein L7a	Hs.99858.0	gb:NM_000972.1	RPL7A	9	3.60	0.0441
179	200801_x_at	actin, beta	Hs.288061.0	gb:NM_001101.2	ACTB	7	3.65	0.0441
180	201077_s_at	non-histone chromosome protein 2 (S. cerevisiae)-like 1	Hs.182255.0	gb:AF155235.1		22	2.72	0.0442
181	217398_x_at	glyceraldehyde-3-phosphate dehydrogenase	Hs.169476.5	gb:AK026525.1		12	3.58	0.0442
182	200001_at	calpain 4, small subunit (30K)	Hs.74451.0	gb:NM_001749.1	CAPN4	19	2.91	0.0442
183	211939_x_at	basic transcription factor 3	Hs.101025.2	gb:X74070.1		5	3.12	0.0442
184	204386_s_at	hypothetical protein MGC3243	Hs.182695.0	gb:BF303597		13	2.68	0.0442
185	212716_s_at	LATS (large tumor suppressor, Drosophila) homolog 1	Hs.324925.2	gb:AW083133_RC		19	2.79	0.0443
186	208114_s_at			gb:NM_030980.1	FLJ12671		2.09	0.0443

Fig. 7

	A	B	C	D	E	F	G	H
187	211784_s_at			gb:BC006181.1			2.39	0.0443
188	221891_x_at	heat shock 70kD protein 8	Hs.180414.3	gb:AA704004_RC		19	3.05	0.0444
189	212790_x_at	ribosomal protein L13a	Hs.119122.2	gb:BF942308_RC		13	3.65	0.0444
190	203012_x_at	ribosomal protein L23a	Hs.184776.0	gb:NM_000984.1	RPL23A	3	3.60	0.0444
191	218226_s_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15)	Hs.227750.0	gb:NM_004547.2	NDUFB4	3	3.00	0.0445
192	218250_s_at	CCR4-NOT transcription complex, subunit 7	Hs.226318.0	gb:NM_013354.2	CNOT7	8	2.44	0.0445
193	213860_x_at	acidic protein rich in leucines	Hs.84264.2	gb:AW268585_RC		5	2.63	0.0446
194	218955_at	hypothetical protein FLJ11052	Hs.274136.0	gb:NM_018310.1	FLJ11052	8	2.18	0.0447
195	200735_x_at	nascent-polypeptide-associated complex alpha polypeptide	Hs.32916.0	gb:NM_005594.1	NACA	12	3.24	0.0447
196	210927_x_at	Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRN.	Hs.323093.0	gb:BC004239.1		1	3.07	0.0449
197	209143_s_at	chloride channel, nucleotide-sensitive, 1A	Hs.84974.0	gb:AF005422.1		11	2.22	0.0449
198	213801_x_at	laminin receptor 1 (67kD, ribosomal protein SA)	Hs.181357.2	gb:AW304232_RC		9	3.32	0.0450
199	208755_x_at	H3 histone, family 3A	Hs.181307.2	gb:BF312331		1	3.47	0.0450
200	208855_s_at	serinethreonine kinase 24 (Ste20, yeast homolog)	Hs.168913.1	gb:AF083420.1	STK3	13	2.86	0.0451
201	201049_s_at	ribosomal protein S18	Hs.275865.0	gb:NM_022551.1	RPS18	6	3.61	0.0451
202	210968_s_at	reticulon 4	Hs.65450.2	gb:AF333336.1		2	3.06	0.0452
203	201544_x_at	poly(A)-binding protein, nuclear 1	Hs.117176.0	gb:BF675004		14	2.75	0.0452
204	200095_x_at	ribosomal protein S10	Hs.76230.2	gb:AA320764_RC		11	3.64	0.0452
205	212933_x_at	ribosomal protein L13	Hs.180842.1	gb:AA961748_RC		16	3.31	0.0452
206	200025_s_at	ribosomal protein L27	Hs.111611.0	gb:NM_000988.1	RPL27	17	3.39	0.0453
207	200674_s_at	ribosomal protein L32	Hs.169793.0	gb:NM_000994.1	RPL32	3	3.50	0.0453
208	211662_s_at			gb:L08666.1	por; por	10	2.96	0.0454
209	208705_s_at	eukaryotic translation initiation factor 5	Hs.286236.1	gb:BG481972		14	2.75	0.0454
210	201994_at	MORF-related gene X	Hs.173714.0	gb:NM_012286.1	KIAA002	X	3.03	0.0455
211	201453_x_at	Ras homolog enriched in brain 2	Hs.279903.0	gb:NM_005614.1	RHEB2	7	2.87	0.0456
212	208980_s_at	ubiquitin C	Hs.183704.1	gb:M26880.1		7	3.33	0.0456
213	200057_s_at	non-POU-domain-containing, octamer-binding	Hs.172207.0	gb:NM_007363.2	NONO	X	2.97	0.0456
214	217719_at	HSPC025	Hs.119503.0	gb:NM_016091.1	HSPC025	22	3.06	0.0458
215	55705_at						2.85	0.0459
216	200826_at	small nuclear ribonucleoprotein D2 polypeptide (16.5kD)	Hs.53125.0	gb:NM_004597.3	SNRPD2	19	2.78	0.0459
217	200064_at	heat shock 90kD protein 1, beta	Hs.74335.0	gb:AF275719.1	HSP90BE	6	3.21	0.0459

Fig. 8

	A	B	C	D	E	F	G	H
218	202836_s_at	similar to S. pombe dim1+	Hs.5074.0	gb:NM_006701.1	DIM1	18	2.63	0.0459
219	206621_s_at	Williams-Beuren syndrome chromosome region 1	Hs.180900.0	gb:NM_022170.1	WBSCR1	7	2.76	0.0460
220	211678_s_at			gb:AF090934.1			2.33	0.0460
221	210501_x_at	LATS (large tumor suppressor, Drosophila) homolog 1	Hs.324925.1	gb:AF119846.1		19	2.84	0.0461
222	209001_s_at	DKFZP566D193 protein	Hs.106909.0	gb:BC005398.1		3	2.76	0.0461
223	212519_at	ubiquitin-conjugating enzyme E2E 1 (homologous to yeast UBC45)	Hs.7766.0	gb:AL518159		3	2.51	0.0461
224	200062_s_at	ribosomal protein L30	Hs.111222.0	gb:L05095.1		8	3.52	0.0461
225	201788_at	RNA helicase-related protein	Hs.8765.0	gb:NM_007372.1	RNAHP	17	2.36	0.0462
226	200086_s_at	cytochrome c oxidase subunit IV	Hs.113205.2	gb:AA854966_RC		16	2.87	0.0462
227	200933_x_at	ribosomal protein S4, X-linked	Hs.108124.0	gb:NM_001007.1	RPS4X	X	3.51	0.0462
228	213828_x_at	H3 histone, family 3A	Hs.181307.4	gb:AA477655		2	3.48	0.0462
229	214143_x_at	ribosomal protein L24	Hs.184582.1	gb:AI560573_RC		9	3.33	0.0462
230	200017_at	ribosomal protein S27a	Hs.3297.0	gb:NM_002954.1	RPS27A	2	3.31	0.0463
231	208909_at	ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1	Hs.3712.0	gb:BC000649.1		19	3.01	0.0463
232	208645_s_at	ribosomal protein S14	Hs.244621.0	gb:AF116710.1		12	3.50	0.0463
233	210434_x_at	Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRN	Hs.323093.1	gb:AF151056.1		1	3.07	0.0464
234	200726_at	protein phosphatase 1, catalytic subunit, gamma isoform	Hs.79081.0	gb:NM_002710.1	PPP1CC	12	2.76	0.0466
235	217769_s_at	hypothetical protein	Hs.279813.0	gb:NM_015932.1	HSPC014	13	2.79	0.0467
236	200775_s_at	heterogeneous nuclear ribonucleoprotein K	Hs.129548.0	gb:BC000355.1		9	2.85	0.0467
237	211058_x_at			gb:BC006379.1		12	3.52	0.0467
238	200876_s_at	proteasome (prosome, macropain) subunit, beta type, 1	Hs.75748.0	gb:NM_002793.1	PSMB1	6	2.59	0.0467
239	201433_s_at	phosphatidylserine synthase 1	Hs.77329.0	gb:NM_014754.1	PTDSS1	8	2.62	0.0467
240	202121_s_at	putative breast adenocarcinoma marker (32kD)	Hs.12107.0	gb:NM_014453.1	BC-2	NA	2.37	0.0470
241	205788_s_at	KIAA0663 gene product	Hs.17969.0	gb:NM_014827.1	KIAA066	6	2.74	0.0471
242	208870_x_at	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polype	Hs.155433.1	gb:BC000931.2		10	3.03	0.0471
243	202243_s_at	proteasome (prosome, macropain) subunit, beta type, 4	Hs.89545.0	gb:NM_002796.1	PSMB4	1	2.54	0.0472
244	201672_s_at	ubiquitin specific protease 14 (tRNA-guanine transglycosylase)	Hs.75981.0	gb:NM_005151.1	USP14	18	2.57	0.0473
245	208635_x_at	ATP-binding cassette, sub-family B (MDRTAP), member 2	Hs.158164.1	gb:BF976260		12	3.29	0.0473
246	201653_at	cornichon-like	Hs.201673.0	gb:NM_005776.1	CNIL	14	2.48	0.0474
247	200794_x_at	DAZ associated protein 2	Hs.75416.0	gb:NM_014764.1	DAZAP2	2	2.89	0.0474
248	201698_s_at	splicing factor, arginineserine-rich 9	Hs.77608.0	gb:NM_003769.1	SFRS9	12	2.64	0.0474

Fig. 9

	A	B	C	D	E	F	G	H
249	208643_s_at	X-ray repair complementing defective repair in Chinese hamster cells 5 (do	Hs.84981.0	gb:J04977.1	G22P1	2	2.58	0.0475
250	202713_s_at	KIAA0391 gene product	Hs.154668.0	gb:AA129755_RC		14	2.16	0.0475
251	210532_s_at	chromosome 14 open reading frame 2	Hs.109052.1	gb:AF116639.1		14	3.00	0.0477
252	200949_x_at	ribosomal protein S20	Hs.8102.0	gb:NM_001023.1	RPS20	8	3.32	0.0477
253	214629_x_at	reticulon 4	Hs.65450.7	gb:AF320999.1		2	3.03	0.0478
254	208680_at	peroxiredoxin 1	Hs.180909.0	gb:L19184.1	NKEFA	1	3.09	0.0478
255	209224_s_at	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)	Hs.163867.0	gb:BC003674.1		5	2.57	0.0479
256	218101_s_at	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B	Hs.193313.0	gb:NM_004549.1	NDUFC2	11	2.90	0.0480
257	213699_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.74405.2	gb:AA854017_RC		10	2.97	0.0480
258	200029_at	ribosomal protein L19	Hs.252723.0	gb:NM_000981.1	RPL19	17	3.36	0.0480
259	221597_s_at	HSPC171 protein	Hs.279593.1	gb:BC003080.1		16	2.31	0.0480
260	200667_at	ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45)	Hs.118797.0	gb:BF448062_RC		4	2.53	0.0481
261	200804_at	testis enhanced gene transcript (BAX inhibitor 1)	Hs.74637.0	gb:NM_003217.1	TEGT	12	2.92	0.0481
262	222230_s_at	uncharacterized hypothalamus protein HARP11	Hs.274369.1	gb:AK022248.1			2.61	0.0482
263	217812_at	high-glucose-regulated protein 8	Hs.20993.0	gb:NM_016258.1	HGRG8	1	2.35	0.0482
264	200817_x_at	ribosomal protein S10	Hs.76230.0	gb:NM_001014.1	RPS10	6	3.65	0.0483
265	200818_at	ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (olig	Hs.76572.0	gb:NM_001697.1	ATP5O	21	2.96	0.0483
266	201934_at	hypothetical protein PRO2730	Hs.194110.0	gb:N92524_RC		4	2.72	0.0484
267	212600_s_at	ubiquinol-cytochrome c reductase core protein II	Hs.173554.1	gb:AV727381		6	2.65	0.0484
268	217964_at	hypothetical protein FLJ20343	Hs.252692.0	gb:NM_017775.1	FLJ20343	17	2.49	0.0484
269	209609_s_at	Homo sapiens, Similar to RIKEN cDNA C330013D18 gene, clone MGC:1	Hs.288936.0	gb:BC004517.1		1	2.16	0.0484
270	212363_x_at	actin, gamma 1	Hs.14376.7	gb:AU145192_RC		6	3.48	0.0485
271	212237_at	KIAA0978 protein	Hs.3686.0	gb:N64780_RC		20	2.25	0.0485
272	200063_s_at	nucleophosmin (nucleolar phosphoprotein B23, numatrin)	Hs.9614.0	gb:BC002398.1		5	3.09	0.0486
273	209860_s_at	annexin A7	Hs.78637.1	gb:J04543.1		10	2.53	0.0488
274	209076_s_at	hypothetical protein 628	Hs.181349.0	gb:BC000974.2		17	2.58	0.0489
275	218034_at	CGI-135 protein	Hs.84344.0	gb:NM_016068.1	LOC5102	7	2.64	0.0490
276	202829_s_at	synaptobrevin-like 1	Hs.24167.0	gb:NM_005638.1	SYBL1	X	2.30	0.0490
277	200047_s_at	YY1 transcription factor	Hs.97496.0	gb:NM_003403.2	YY1	14	2.61	0.0490
278	200877_at	chaperonin containing TCP1, subunit 4 (delta)	Hs.79150.0	gb:NM_006430.1	CCT4	2	2.94	0.0490
279	200910_at	chaperonin containing TCP1, subunit 3 (gamma)	Hs.1708.0	gb:NM_005998.1	CCT3	1	2.59	0.0491

Fig. 10

	A	B	C	D	E	F	G	H
280	213377_x_at	ribosomal protein S12	Hs.285405.2	gb:AI799007_RC		11	3.48	0.0491
281	214112_s_at	Homo sapiens (clone 115392) mRNA	Hs.82171.4	gb:AA543076_RC		X	2.26	0.0492
282	219035_s_at	hypothetical protein FLJ21786	Hs.316809.0	gb:NM_025126.1	FLJ21786	12	2.35	0.0493
283	218011_at	ubiquitin-like 5	Hs.109701.0	gb:NM_024292.1	UBL5	19	2.31	0.0493
284	201119_s_at	cytochrome c oxidase subunit VIII	Hs.81097.0	gb:NM_004074.1	COX8	11	3.07	0.0495
285	211072_x_at			gb:BC006481.1		12	3.47	0.0496
286	208695_s_at	ribosomal protein L39	Hs.300141.1	gb:BC001019.1		X	3.48	0.0497
287	217742_s_at	hypothetical protein	Hs.70333.0	gb:NM_016628.1	LOC5132	10	2.62	0.0498
288	218191_s_at	hypothetical protein FLJ11240	Hs.301308.0	gb:NM_018368.1	FLJ11240	6	2.77	0.0498
289	211940_x_at	H3 histone, family 3A	Hs.181307.5	gb:BE869922		2	3.42	0.0498
290	201059_at	ems1 sequence (mammary tumor and squamous cell carcinoma-associated (	Hs.119257.0	gb:NM_005231.1	EMS1	11	2.49	0.0499
291	205292_s_at	heterogeneous nuclear ribonucleoprotein A2B1	Hs.75598.0	gb:NM_002137.1	HNRPA2I	7	2.86	0.0499
292	200000_s_at	U5 snRNP-specific protein (220 kD), ortholog of S. cerevisiae Prp8p	Hs.181368.0	gb:NM_006445.1	PRP8	17	2.54	0.0499
293	212137_at	pregnancy-associated plasma protein A	Hs.75874.1	gb:AV746402		5	2.67	0.0500
294	211475_s_at	BCL2-associated athanogene	Hs.41714.1	gb:AF116273.1		9	2.43	0.0500
295	212039_x_at	ribosomal protein L3	Hs.119598.2	gb:BG339228		14	3.59	0.0500
296	208652_at	protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform	Hs.91773.0	gb:BC000400.1		5	2.57	0.0500
297	209329_x_at	mannan-binding lectin serine protease 1 (C4C2 activating component of Ra	Hs.227152.1	gb:BC000587.1		5	2.73	0.0500
298	209452_s_at	vesicle-associated soluble NSF attachment protein receptor (v-SNARE; hor	Hs.169206.0	gb:AF035824.1	VTI1	14	2.53	0.0500
299	221700_s_at			gb:AF348700.1	UBA52	19	3.23	0.0501
300	208687_x_at	heat shock 70kD protein 8	Hs.180414.0	gb:AF352832.1	HSC70	11	3.11	0.0502
301	200748_s_at	ferritin, heavy polypeptide 1	Hs.62954.0	gb:NM_002032.1	FTH1	11	3.32	0.0502
302	200031_s_at	ribosomal protein S11	Hs.182740.0	gb:NM_001015.1	RPS11	19	3.55	0.0503
303	209058_at	endothelial differentiation-related factor 1	Hs.174050.0	gb:AB002282.1		9	2.53	0.0503
304	201338_x_at	general transcription factor IIIA	Hs.75113.0	gb:NM_002097.1	GTF3A	13	2.75	0.0503
305	222047_s_at	arsenate resistance protein ARS2	Hs.111801.2	gb:AI523895_RC		7	2.61	0.0503
306	200871_s_at	prosaposin (variant Gaucher disease and variant metachromatic leukodystro	Hs.78575.0	gb:NM_002778.1	PSAP	10	3.28	0.0504
307	200033_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 5 (RNA helicase, 68kD)	Hs.76053.0	gb:NM_004396.2	DDX5	17	3.00	0.0504
308	218381_s_at	U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	Hs.7655.0	gb:NM_007279.1	U2AF65	19	2.22	0.0505
309	200738_s_at	phosphoglycerate kinase 1	Hs.78771.0	gb:NM_000291.1	PGK1	X	2.94	0.0506
310	214214_s_at	complement component 1, q subcomponent binding protein	Hs.78614.3	gb:AU151801_RC		17	2.23	0.0506

Fig. 11

	A	B	C	D	E	F	G	H
311	208865_at	hypothetical protein PRO2975	Hs.144477.0	gb:BG534245		5	2.68	0.0508
312	201097_s_at	ADP-ribosylation factor 4	Hs.75290.0	gb:NM_001660.2	ARF4	3	2.91	0.0509
313	200716_x_at	ribosomal protein L13a	Hs.119122.0	gb:NM_012423.1	RPL13A	19	3.63	0.0509
314	218357_s_at	translocase of inner mitochondrial membrane 8 (yeast) homolog B	Hs.279915.0	gb:NM_012459.1	TIMM8B	11	2.71	0.0510
315	213545_x_at	sorting nexin 3	Hs.12102.2	gb:BE962615_RC		6	2.76	0.0510
316	202164_s_at	CCR4-NOT transcription complex, subunit 8	Hs.26703.0	gb:AF180476.1	CALIF	5	2.18	0.0511
317	207168_s_at	H2A histone family, member Y	Hs.75258.0	gb:NM_004893.1	H2AFY	5	2.77	0.0511
318	208656_s_at	cyclin I	Hs.79933.0	gb:AF135162.1	CYC1	4	3.15	0.0511
319	210149_s_at	Homo sapiens clone 023e08 My032 protein mRNA, complete cds	Hs.307099.0	gb:AF061735.1		17	3.02	0.0512
320	201254_x_at	ribosomal protein S6	Hs.241507.0	gb:NM_001010.1	RPS6	9	3.55	0.0512
321	201600_at	B-cell associated protein	Hs.7771.0	gb:NM_007273.1	REA	12	3.01	0.0512
322	215383_x_at	Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K	Hs.293681.0	gb:AL137312.1		15	2.24	0.0513
323	205711_x_at	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypep	Hs.155433.0	gb:NM_005174.1	ATP5C1	10	2.94	0.0513
324	213366_x_at	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypep	Hs.155433.3	gb:AV711183		14	3.00	0.0513
325	212199_at	Human putative ribosomal protein S1 mRNA	Hs.284281.0	gb:AL566962_RC		4	2.31	0.0513
326	200034_s_at	ribosomal protein L6	Hs.174131.0	gb:NM_000970.1	RPL6	12	3.15	0.0514
327	200014_s_at	heterogeneous nuclear ribonucleoprotein C (C1C2)	Hs.182447.0	gb:NM_004500.1	HNRPC	14	2.47	0.0515
328	203107_x_at	ribosomal protein S2	Hs.182426.0	gb:NM_002952.1	RPS2	16	3.62	0.0516
329	202220_at	KIAA0907 protein	Hs.24656.0	gb:NM_014949.1	KIAA090	1	2.37	0.0516
330	207507_s_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subu	Hs.429.0	gb:NM_001689.1	ATP5G3	2	2.88	0.0516
331	210949_s_at	eukaryotic translation initiation factor 3, subunit 8 (110kD)	Hs.4835.1	gb:BC000533.1		16	2.96	0.0516
332	219293_s_at	hypothetical protein	Hs.86347.0	gb:NM_013341.1	PTD004	22	2.59	0.0516
333	201597_at	cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	Hs.70312.0	gb:NM_001865.1	COX7A2	6	3.01	0.0516
334	221434_s_at			gb:NM_031210.1	DC50	14	2.57	0.0516
335	201145_at	HS1 binding protein	Hs.15318.0	gb:NM_006118.2	HAX1	1	2.54	0.0517
336	200091_s_at	ribosomal protein S25	Hs.113029.1	gb:AA888388_RC		5	3.05	0.0517
337	212773_s_at	translocase of outer mitochondrial membrane 20 (yeast) homolog	Hs.75187.2	gb:BG165094		7	2.41	0.0517
338	203445_s_at	conserved gene amplified in osteosarcoma	Hs.180669.0	gb:NM_005730.1	OS4	6	2.81	0.0518
339	200693_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.74405.0	gb:NM_006826.1	YWHAQ	2	3.00	0.0519
340	200714_x_at	amplified in osteosarcoma	Hs.76228.0	gb:NM_006812.1	OS-9	12	2.82	0.0519
341	213941_x_at	ribosomal protein S7	Hs.301547.3	gb:AI970731_RC		19	3.21	0.0519

Fig. 12

	A	B	C	D	E	F	G	H
342	217906_at	host cell factor homolog	Hs.20597.0	gb:NM_014315.1	LCP		2.64	0.0519
343	203075_at	MAD (mothers against decapentaplegic, Drosophila) homolog 2	Hs.82483.0	gb:AW151617_RC		18	2.20	0.0519
344	213646_x_at	hypothetical protein FLJ13556 similar to N-myc downstream regulated 3	Hs.240615.2	gb:BE300252_RC		12	3.50	0.0519
345	201217_x_at	ribosomal protein L3	Hs.119598.0	gb:NM_000967.1	RPL3	22	3.59	0.0520
346	213619_at	heterogeneous nuclear ribonucleoprotein H1 (H)	Hs.245710.1	gb:AV753392		1	3.03	0.0521
347	203136_at	Rab acceptor 1 (prenylated)	Hs.11417.0	gb:NM_006423.1	RABAC1	19	2.68	0.0522
348	204839_at	hypothetical protein	Hs.279913.0	gb:NM_015918.1	HSPC004	12	2.26	0.0522
349	211769_x_at			gb:BC006088.1		20	2.76	0.0522
350	215628_x_at	Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M1	Hs.302053.0	gb:AL049285.1		5	2.19	0.0522
351	200022_at	ribosomal protein L18	Hs.75458.0	gb:NM_000979.1	RPL18	19	3.31	0.0523
352	202160_at	CREB binding protein (Rubinstein-Taybi syndrome)	Hs.23598.0	gb:NM_004380.1	CREBBP	16	2.22	0.0523
353	221483_s_at	cyclic AMP phosphoprotein, 19 kD	Hs.7351.0	gb:AF084555.1	ARPP-19	15	2.73	0.0523
354	218334_at	hypothetical protein FLJ23445	Hs.288151.0	gb:NM_025075.1	FLJ23445	3	2.14	0.0523
355	200046_at	defender against cell death 1	Hs.82890.0	gb:NM_001344.1	DAD1	14	2.59	0.0524
356	200027_at	asparaginyl-tRNA synthetase	Hs.181311.0	gb:NM_004539.2	NARS	18	2.81	0.0524
357	200680_x_at	high-mobility group (nonhistone chromosomal) protein 1	Hs.274472.0	gb:NM_002128.1	HMG1	13	3.04	0.0524
358	204640_s_at	speckle-type POZ protein	Hs.129951.0	gb:NM_003563.1	SPOP	17	2.38	0.0524
359	200840_at	lysyl-tRNA synthetase	Hs.3100.0	gb:NM_005548.1	KARS	16	2.82	0.0525
360	203462_x_at	eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)	Hs.57783.0	gb:NM_003751.1	EIF3S9	7	2.54	0.0526
361	200781_s_at	ribosomal protein S15a	Hs.2953.0	gb:NM_001019.1	RPS15A	16	3.36	0.0527
362	200048_s_at	jumping translocation breakpoint	Hs.6396.0	gb:NM_006694.1	JTB	1	2.95	0.0527
363	200640_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.75103.0	gb:NM_003406.1	YWHAZ	8	2.89	0.0527
364	201757_at	NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) (NADH-coenzy	Hs.80595.0	gb:NM_004552.1	NDUFS5	1	2.97	0.0527
365	213080_x_at	ribosomal protein L5	Hs.180946.3	gb:BF214492		1	3.30	0.0528
366	213702_x_at	N-acylsphingosine amidohydrolase (acid ceramidase)	Hs.75811.3	gb:AI934569_RC		8	3.04	0.0528
367	213513_x_at	Rho guanine nucleotide exchange factor (GEF) 1	Hs.252280.1	gb:BG034239		2	2.67	0.0529
368	200763_s_at	ribosomal protein, large, P1	Hs.177592.0	gb:NM_001003.1	RPLP1	15	3.51	0.0529
369	212787_at	ZAP3 protein	Hs.159471.0	gb:AI952986_RC		14	2.27	0.0529
370	202077_at	NADH dehydrogenase (ubiquinone) 1, alphabeta subcomplex, 1 (8kD, SDA	Hs.5556.0	gb:NM_005003.1	NDUFAB	16	2.87	0.0529
371	201239_s_at	KIAA0102 gene product	Hs.77665.0	gb:BF530535		11	2.55	0.0530
372	217917_s_at	HSPC162 protein	Hs.100002.0	gb:NM_014183.1	HSPC162	20	2.78	0.0530

Fig. 13

	A	B	C	D	E	F	G	H
373	40446_at						2.68	0.0530
374	200927_s_at	GTPase Rab14	Hs.5807.0	gb:AA919115_RC		9	2.27	0.0530
375	200089_s_at	ribosomal protein L4	Hs.286.4	gb:AI953886_RC		9	3.10	0.0531
376	201092_at	retinoblastoma-binding protein 7	Hs.31314.0	gb:NM_002893.2	RBBP7	X	2.64	0.0531
377	200099_s_at	Human DNA sequence from clone RP11-486O22 on chromosome 10 Cont	Hs.307132.0	gb:AL356115			3.40	0.0531
378	206790_s_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL)	Hs.183435.0	gb:NM_004545.1	NDUFB1	14	2.87	0.0532
379	211509_s_at	reticulon 4	Hs.65450.4	gb:AB015639.1	ASY	2	3.10	0.0532
380	207812_s_at	DKFZP434D156 protein	Hs.6880.0	gb:NM_015530.1	DKFZP43	2	2.39	0.0533
381	209075_s_at	nitrogen fixation cluster-like	Hs.9908.1	gb:AY009128.1	ISCU	12	2.91	0.0533
382	201901_s_at	YY1 transcription factor	Hs.97496.0	gb:Z14077.1		14	2.51	0.0533
383	200013_at	ribosomal protein L24	Hs.184582.0	gb:NM_000986.1	RPL24	3	3.22	0.0534
384	37012_at						2.81	0.0534
385	218059_at	HSPC038 protein	Hs.23528.0	gb:NM_016096.1	LOC5112	8	2.62	0.0534
386	208752_x_at	nucleosome assembly protein 1-like 1	Hs.179662.1	gb:AI888672_RC		12	2.84	0.0534
387	218213_s_at	chromosome 11 open reading frame 10	Hs.90918.0	gb:NM_014206.1	C11orf10	11	2.93	0.0535
388	201891_s_at	beta-2-microglobulin	Hs.75415.0	gb:NM_004048.1	B2M	15	3.44	0.0535
389	212352_s_at	transmembrane trafficking protein	Hs.74137.2	gb:BE780075		14	2.77	0.0535
390	200002_at	ribosomal protein L35	Hs.182825.0	gb:NM_007209.1	RPL35	9	3.20	0.0536
391	208620_at	poly(rC)-binding protein 1	Hs.2853.0	gb:U24223.1	alpha-CP1	2	2.96	0.0536
392	222021_x_at	succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	Hs.469.2	gb:AI348006_RC		3	2.70	0.0536
393	213890_x_at	ribosomal protein S16	Hs.80617.1	gb:AI200589_RC		19	3.43	0.0536
394	200705_s_at	eukaryotic translation elongation factor 1 beta 2	Hs.275959.0	gb:NM_001959.1	EEF1B2	2	3.14	0.0537
395	213347_x_at	ribosomal protein S4, X-linked	Hs.108124.5	gb:AW132023_RC		10	3.36	0.0537
396	212802_s_at	DKFZP434C212 protein	Hs.172069.0	gb:AK023841.1		9	2.16	0.0537
397	221488_s_at	divalent cation tolerant protein CUTA	Hs.107187.1	gb:AF230924.1		6	2.78	0.0537
398	203316_s_at	small nuclear ribonucleoprotein polypeptide E	Hs.1066.0	gb:NM_003094.1	SNRPE	1	2.53	0.0539
399	200822_x_at	triosephosphate isomerase 1	Hs.83848.0	gb:NM_000365.1	TPI1	12	3.04	0.0539
400	208768_x_at	ribosomal protein L22	Hs.99914.1	gb:D17652.1		1	3.33	0.0539
401	200709_at	FK506-binding protein 1A (12kD)	Hs.752.0	gb:NM_000801.1	FKBP1A	20	2.79	0.0539
402	200009_at	GDP dissociation inhibitor 2	Hs.56845.0	gb:NM_001494.2	GDI2	10	2.78	0.0539
403	201343_at	ubiquitin-conjugating enzyme E2D 2 (homologous to yeast UBC45)	Hs.108332.0	gb:BE621259		5	2.49	0.0540

Fig. 14

	A	B	C	D	E	F	G	H
404	211927_x_at	eukaryotic translation elongation factor 1 gamma	Hs.2186.3	gb:BE963164_RC		7	3.44	0.0541
405	213360_s_at	KIAA0618 gene product	Hs.295112.4	gb:AA514622_RC		7	2.67	0.0541
406	209004_s_at	f-box and leucine-rich repeat protein 5	Hs.5548.0	gb:AF142481.1	FLR1	4	2.48	0.0541
407	200038_s_at	ribosomal protein L17	Hs.82202.0	gb:NM_000985.1	RPL17	18	3.28	0.0542
408	218106_s_at	hypothetical protein FLJ10567	Hs.28444.0	gb:NM_018141.1	FLJ10567	6	2.08	0.0543
409	200055_at	TATA box binding protein (TBP)-associated factor, RNA polymerase II, H	Hs.89657.0	gb:NM_006284.1	TAF2H	11	2.70	0.0544
410	211073_x_at			gb:BC006483.1		22	3.56	0.0544
411	35150_at						2.40	0.0544
412	208833_s_at	like mouse brain protein E46	Hs.13493.0	gb:AF119662.1		22	2.89	0.0544
413	200079_s_at	lysyl-tRNA synthetase	Hs.3100.1	gb:AF285758.1		16	2.61	0.0546
414	202325_s_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6	Hs.73851.0	gb:NM_001685.1	ATP5J	21	2.74	0.0546
415	217724_at	PAI-1 mRNA-binding protein	Hs.165998.0	gb:AF131807.1		1	2.84	0.0546
416	200010_at	ribosomal protein L11	Hs.179943.0	gb:NM_000975.1	RPL11	1	3.27	0.0546
417	209089_at	RAB5A, member RAS oncogene family	Hs.73957.1	gb:BC001267.1		4	2.59	0.0547
418	208852_s_at	calnexin	Hs.155560.0	gb:AI761759_RC		UL	2.63	0.0547
419	200026_at	ribosomal protein L34	Hs.250895.0	gb:NM_000995.1	RPL34	4	3.39	0.0547
420	203221_at	transducin-like enhancer of split 1, homolog of Drosophila E(sp1)	Hs.28935.0	gb:AI758763_RC		9	2.38	0.0547
421	200614_at	clathrin, heavy polypeptide (Hc)	Hs.178710.0	gb:NM_004859.1	CLTC	17	3.05	0.0547
422	213101_s_at	hypothetical protein EUROIMAGE1875335	Hs.10927.1	gb:Z78330		2	3.00	0.0548
423	200779_at	activating transcription factor 4 (tax-responsive enhancer element B67)	Hs.181243.0	gb:NM_001675.1	ATF4	22	3.00	0.0549
424	200964_at	ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature sensitivity	Hs.2055.0	gb:NM_003334.1	UBE1		2.67	0.0549
425	201094_at	ribosomal protein S29	Hs.539.0	gb:NM_001032.1	RPS29	14	3.42	0.0550
426	215952_s_at	ornithine decarboxylase antizyme 1	Hs.125078.2	gb:AF090094.1		19	2.98	0.0551
427	215091_s_at	general transcription factor IIIA	Hs.75113.3	gb:BE542815		13	2.68	0.0551
428	200754_x_at	splicing factor, arginineserine-rich 2	Hs.73965.0	gb:NM_003016.1	SFRS2	17	2.84	0.0551
429	207040_s_at	suppression of tumorigenicity 13 (colon carcinoma) (Hsp70-interacting pro	Hs.119222.0	gb:NM_003932.1	ST13	7	2.88	0.0552
430	217861_s_at	prolactin regulatory element binding	Hs.279784.0	gb:NM_013388.1	PREB	2	2.17	0.0552
431	207508_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subu	Hs.429.0	gb:NM_001689.1	ATP5G3	2	2.84	0.0553
432	216396_s_at	etoposide-induced mRNA	Hs.286027.3	gb:AF131850.1		11	2.28	0.0553
433	208655_at	cyclin I	Hs.79933.0	gb:BG530368		4	2.95	0.0554
434	201029_s_at	antigen identified by monoclonal antibodies 12E7, F21 and O13	Hs.177543.0	gb:NM_002414.1	MIC2	X	3.00	0.0554

Fig. 15

	A	B	C	D	E	F	G	H
435	201019_s_at	eukaryotic translation initiation factor 1A	Hs.4310.0	gb:NM_001412.1	EIF1A	1	2.50	0.0554
436	211345_x_at	eukaryotic translation elongation factor 1 gamma	Hs.2186.1	gb:AF119850.1			3.40	0.0555
437	212867_at	nuclear receptor coactivator 2	Hs.29131.1	gb:AI040324_RC		8	2.12	0.0556
438	201322_at	ATP synthase, H+ transporting, mitochondrial F1 complex, beta polypeptid	Hs.25.0	gb:NM_001686.1	ATP5B	12	3.15	0.0556
439	221775_x_at	ribosomal protein L22	Hs.99914.2	gb:BG152979		3	3.28	0.0556
440	209134_s_at	ribosomal protein S6	Hs.241507.1	gb:BC000524.1		19	3.48	0.0556
441	213588_x_at	ribosomal protein L14	Hs.738.4	gb:AA838274_RC		3	3.05	0.0556
442	200958_s_at	syndecan binding protein (syntenin)	Hs.8180.0	gb:NM_005625.1	SDCBP	8	2.83	0.0556
443	212071_s_at	Homo sapiens cDNA FLJ13700 fis, clone PLACE2000216, highly similar t	Hs.324648.0	gb:BE968833		2	3.08	0.0559
444	213404_s_at	Ras homolog enriched in brain 2	Hs.279903.2	gb:BF033683_RC		7	2.51	0.0559
445	200023_s_at	eukaryotic translation initiation factor 3, subunit 5 (epsilon, 47kD)	Hs.7811.0	gb:NM_003754.1	EIF3S5	2	2.90	0.0560
446	203621_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5 (16kD, SGDH)	Hs.19236.0	gb:NM_002492.1	NDUFB5	3	2.65	0.0560
447	200675_at	CD81 antigen (target of antiproliferative antibody 1)	Hs.54457.0	gb:NM_004356.1	CD81	11	3.13	0.0561
448	50277_at						2.75	0.0561
449	208828_at	histone fold protein CHRAC17; DNA polymerase epsilon p17 subunit	Hs.108112.0	gb:BC004170.1		9	2.15	0.0565
450	210275_s_at	zinc finger protein 216	Hs.3776.1	gb:AF062347.1	ZNF216	9	2.95	0.0566
451	221471_at	tumor differentially expressed 1	Hs.272168.0	gb:AW173623_RC		20	2.50	0.0566
452	201154_x_at	ribosomal protein L4	Hs.286.0	gb:NM_000968.1	RPL4	3	3.29	0.0566
453	218165_at	hypothetical protein FLJ11730	Hs.17118.0	gb:NM_022756.1	FLJ11730	1	2.30	0.0567
454	214113_s_at	RNA binding motif protein 8A	Hs.65648.1	gb:AI738479_RC		14	1.90	0.0567
455	201031_s_at	heterogeneous nuclear ribonucleoprotein H1 (H)	Hs.245710.0	gb:NM_005520.1	HNRPH1	1	2.67	0.0567
456	201899_s_at	ubiquitin-conjugating enzyme E2A (RAD6 homolog)	Hs.80612.0	gb:NM_003336.1	UBE2A	X	2.36	0.0567
457	217938_s_at	potassium channel modulatory factor	Hs.5392.0	gb:NM_020122.1	DKFZP43	2	2.57	0.0568
458	212461_at	protease, serine, 15	Hs.278614.1	gb:BF793951		8	2.50	0.0568
459	218633_x_at	hypothetical protein FLJ11342	Hs.266514.0	gb:NM_018394.1	FLJ11342	3	2.41	0.0569
460	208737_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump), member J	Hs.90336.0	gb:BC003564.1		9	2.71	0.0570
461	211487_x_at	ribosomal protein S17	Hs.5174.1	gb:BC004886.1			3.49	0.0571
462	214259_s_at	aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)	Hs.6980.1	gb:AI144075			2.33	0.0571
463	211710_x_at			gb:BC005817.1		3	3.32	0.0574
464	208706_s_at	eukaryotic translation initiation factor 5	Hs.286236.1	gb:AK026933.1		14	2.49	0.0574
465	209141_at	ubiquitin-conjugating enzyme E2G 1 (homologous to C. elegans UBC7)	Hs.78563.0	gb:AW299555_RC		17	2.46	0.0575

Fig. 16

	A	B	C	D	E	F	G	H
466	202591_s_at	single-stranded DNA-binding protein	Hs.923.0	gb:NM_003143.1	SSBP	7	2.34	0.0575
467	212041_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump), member D	Hs.106876.0	gb:AL566172_RC		16	2.72	0.0576
468	200937_s_at	ribosomal protein L5	Hs.180946.0	gb:NM_000969.1	RPL5	1	3.17	0.0577
469	201132_at	heterogeneous nuclear ribonucleoprotein H2 (H)	Hs.278857.0	gb:NM_019597.1	HNRPH2	X	2.20	0.0578
470	201517_at	nuclear cap binding protein subunit 2, 20kD	Hs.240770.0	gb:BC001255.1		3	2.09	0.0578
471	210097_s_at	retinoic acid repressible protein	Hs.106346.1	gb:AF130102.1_RC		6	2.34	0.0578
472	200037_s_at	heterochromatin-like protein 1	Hs.278554.0	gb:NM_016587.1	HECH	7	2.51	0.0578
473	45572_s_at						2.71	0.0579
474	212038_s_at	voltage-dependent anion channel 1	Hs.149155.2	gb:AL515918_RC		5	2.60	0.0580
475	200774_at	C9orf10 protein	Hs.76666.0	gb:BE963765_RC		9	2.74	0.0580
476	217802_s_at	similar to rat nuclear ubiquitous casein kinase 2	Hs.118064.0	gb:NM_022731.1	NUCKS	1	2.59	0.0581
477	220856_x_at	PRO0470 protein	Hs.278937.0	gb:NM_014128.1	PRO0470		2.67	0.0582
478	213687_s_at	uncharacterized hypothalamus protein HSMNP1	Hs.179666.2	gb:BE968801		12	3.15	0.0582
479	201358_s_at	coatomer protein complex, subunit beta	Hs.3059.0	gb:NM_016451.1	COPB	11	2.70	0.0583
480	200689_x_at	eukaryotic translation elongation factor 1 gamma	Hs.2186.0	gb:NM_001404.1	EEF1G	11	3.37	0.0583
481	40225_at						2.63	0.0584
482	36994_at						3.03	0.0585
483	208638_at	protein disulfide isomerase-related protein	Hs.182429.1	gb:BE910010		2	2.44	0.0585
484	212639_x_at	Tubulin, alpha, brain-specific	Hs.272897.2	gb:AL581768_RC		12	3.45	0.0586
485	208764_s_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subu	Hs.89399.1	gb:D13119.1	P2	12	2.66	0.0586
486	209377_s_at	thyroid hormone receptor interactor 7	Hs.77558.0	gb:AF274949.1		6	2.81	0.0586
487	217707_x_at	ESTs	Hs.327729.0	gb:AI535683_RC		9	2.36	0.0587
488	213735_s_at	cytochrome c oxidase subunit Vb	Hs.1342.1	gb:AI557312		2	3.07	0.0587
489	201078_at	transmembrane 9 superfamily member 2	Hs.28757.0	gb:NM_004800.1	TM9SF2	13	2.71	0.0587
490	221619_s_at	mitochondrial carrier homolog 1	Hs.279939.1	gb:AF189289.1		6	3.13	0.0589
491	201281_at	cell membrane glycoprotein, 110000M(r) (surface antigen)	Hs.90107.0	gb:NM_007002.1	GP110	20	2.36	0.0589
492	202233_s_at	ubiquinol-cytochrome c reductase hinge protein	Hs.73818.0	gb:NM_006004.1	UQCRH	1	2.86	0.0590
493	208924_at	ring finger protein 11	Hs.96334.0	gb:AB024703.1	Sid1669	1	2.63	0.0590
494	213086_s_at	acidic protein rich in leucines	Hs.84264.3	gb:BF341845			2.40	0.0591
495	201068_s_at	proteasome (prosome, macropain) 26S subunit, ATPase, 2	Hs.61153.0	gb:NM_002803.1	PSMC2	3	2.56	0.0592
496	215963_x_at	Human DNA sequence from clone RP1-111B22 on chromosome 6q16-21 C	Hs.296559.0	gb:Z98200		6	3.48	0.0592

Fig. 17

	A	B	C	D	E	F	G	H
497	209492_x_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e	Hs.85539.1	gb:BC003679.1			2.75	0.0592
498	212967_x_at	nucleosome assembly protein 1-like 1	Hs.179662.2	gb:AW148801_RC		12	2.82	0.0593
499	40189_at						2.53	0.0593
500	221743_at	DnaJ (Hsp40) homolog, subfamily C, member 3	Hs.9683.1	gb:AI472139_RC		11	2.61	0.0595
501	212622_at	KIAA0033 protein	Hs.174905.0	gb:N64760_RC		2	2.26	0.0595
502	200892_s_at	splicing factor, arginineserine-rich (transformer 2 Drosophila homolog) 10	Hs.30035.0	gb:BC000451.1		3	2.35	0.0596
503	217827_s_at	acid cluster protein 33	Hs.71475.0	gb:NM_016630.2	ACP33	15	2.47	0.0596
504	201352_at	YME1 (S.cerevisiae)-like 1	Hs.206521.0	gb:NM_014263.1	YME1L1	10	2.55	0.0598
505	201256_at	cytochrome c oxidase subunit VIIa polypeptide 2 like	Hs.30888.0	gb:NM_004718.1	COX7A2l	2	2.64	0.0598
506	200694_s_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 24	Hs.155986.0	gb:NM_020414.2	DDX24	14	2.69	0.0598
507	201552_at	lysosomal-associated membrane protein 1	Hs.150101.0	gb:NM_005561.2	LAMP1	13	2.80	0.0598
508	211942_x_at	ribosomal protein L13a	Hs.119122.3	gb:BF979419		10	3.38	0.0599
509	221473_x_at	tumor differentially expressed 1	Hs.272168.0	gb:U49188.1	Diff33	20	2.61	0.0600
510	200945_s_at	yeast Sec31p homolog	Hs.70266.1	gb:NM_014933.1	KIAA090	4	2.66	0.0600
511	211997_x_at	H3 histone, family 3B (H3.3B)	Hs.180877.1	gb:NM_005324.1	H3F3B	17	3.26	0.0600
512	220960_x_at	ribosomal protein L22	Hs.99914.0	gb:NM_000983.1	RPL22	1	3.15	0.0600
513	200092_s_at	ribosomal protein L37	Hs.179779.1	gb:BF216701		11	3.27	0.0600
514	221158_at	GC-rich sequence DNA-binding factor candidate	Hs.283771.0	gb:NM_013329.1	GCFC	21	1.86	0.0600
515	60528_at						2.45	0.0601
516	208734_x_at	RAB2, member RAS oncogene family	Hs.78305.0	gb:M28213.1	RAB2	8	2.35	0.0601
517	200854_at	nuclear receptor co-repressor 1	Hs.144904.0	gb:AB028970.1	KIAA104	17	2.13	0.0602
518	208946_s_at	beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)	Hs.12272.0	gb:AF139131.1	BECN1	17	2.20	0.0602
519	209033_s_at	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A	Hs.75842.0	gb:D86550.1	hMNB	21	2.52	0.0603
520	201227_s_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI)	Hs.198273.0	gb:NM_005004.1	NDUFB8	10	2.78	0.0603
521	201366_at	annexin A7	Hs.78637.0	gb:NM_004034.1	ANXA7	10	2.32	0.0604
522	217835_x_at	putative Rab5-interacting protein	Hs.184062.0	gb:NM_018840.1	LOC5596	20	2.82	0.0604
523	208801_at	signal recognition particle 72kD	Hs.237825.0	gb:BE856385_RC		4	2.46	0.0604
524	207628_s_at	putative methyltransferase	Hs.155020.0	gb:NM_017528.1	HASJ444	7	2.28	0.0604
525	218201_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)	Hs.198272.0	gb:NM_004546.1	NDUFB2	7	2.50	0.0606
526	208248_x_at	amyloid beta (A4) precursor-like protein 2	Hs.279518.0	gb:NM_001642.1	APLP2	11	2.84	0.0607
527	212734_x_at	ribosomal protein L13	Hs.180842.4	gb:AI186735_RC		17	3.50	0.0608

Fig. 18

	A	B	C	D	E	F	G	H
528	213318_s_at	HLA-B associated transcript-3	Hs.274348.2	gb:BG028844		6	2.57	0.0608
529	200663_at	CD63 antigen (melanoma 1 antigen)	Hs.76294.0	gb:NM_001780.1	CD63	12	3.20	0.0608
530	202224_at	v-crk avian sarcoma virus CT10 oncogene homolog	Hs.306088.0	gb:BF304695		17	2.15	0.0608
531	217718_s_at	GW128 protein	Hs.182238.0	gb:NM_014052.1	GW128	20	3.00	0.0608
532	201231_s_at	enolase 1, (alpha)	Hs.254105.0	gb:NM_001428.1	ENO1	1	2.95	0.0608
533	218200_s_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)	Hs.198272.0	gb:NM_004546.1	NDUFB2	7	3.00	0.0608
534	200691_s_at	heat shock 70kD protein 9B (mortalin-2)	Hs.3069.0	gb:BC000478.1		5	2.58	0.0609
535	218286_s_at	ring finger protein 7	Hs.14084.0	gb:NM_014245.1	RNF7	3	2.15	0.0609
536	200847_s_at	HSPC035 protein	Hs.279921.0	gb:NM_016127.1	LOC5166	8	2.83	0.0609
537	208843_s_at	DKFZP434D156 protein	Hs.6880.1	gb:BC001408.1		2	2.32	0.0609
538	217736_s_at	heme-regulated initiation factor 2-alpha kinase	Hs.258730.0	gb:NM_014413.2	HRI	7	2.61	0.0609
539	218026_at	HSPC009 protein	Hs.16059.0	gb:NM_014019.1	HSPC009	17	2.69	0.0610
540	213798_s_at	adenylyl cyclase-associated protein	Hs.104125.2	gb:AA806142_RC		1	2.70	0.0611
541	202333_s_at	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	Hs.811.0	gb:AA877765_RC		5	2.42	0.0611
542	202296_s_at	similar to S. cerevisiae RER1	Hs.40500.0	gb:NM_007033.1	RER1	1	2.50	0.0611
543	200650_s_at	lactate dehydrogenase A	Hs.2795.0	gb:NM_005566.1	LDHA	11	3.14	0.0612
544	213612_x_at	hypothetical protein	Hs.218329.3	gb:AI800419_RC		1	2.95	0.0613
545	209115_at	ubiquitin-activating enzyme E1C (homologous to yeast UBA3)	Hs.154320.0	gb:AL117566.1	DKFZp56	3	2.10	0.0615
546	213048_s_at	SET translocation (myeloid leukemia-associated)	Hs.145279.2	gb:W26593		9	2.81	0.0615
547	202343_x_at	cytochrome c oxidase subunit Vb	Hs.1342.0	gb:NM_001862.1	COX5B	2	2.90	0.0616
548	201258_at	ribosomal protein S16	Hs.80617.0	gb:NM_001020.1	RPS16	19	3.25	0.0616
549	217733_s_at	thymosin, beta 10	Hs.76293.0	gb:NM_021103.1	TMSB10	2	3.37	0.0616
550	210774_s_at	nuclear receptor coactivator 4	Hs.99908.0	gb:AL162047.1	DKFZp76	10	2.89	0.0617
551	220746_s_at	retinoid x receptor interacting protein	Hs.7889.0	gb:NM_016290.1	LOC5172	5	2.13	0.0618
552	201807_at	vacuolar protein sorting 26 (yeast homolog)	Hs.67052.0	gb:NM_004896.1	VPS26	10	2.56	0.0619
553	200866_s_at	prosaposin (variant Gaucher disease and variant metachromatic leukodystro	Hs.78575.0	gb:M32221.1	PSAP	10	2.99	0.0619
554	202810_at	developmentally regulated GTP-binding protein 1	Hs.115242.0	gb:NM_004147.1	DRG1	22	2.45	0.0621
555	200896_x_at	hepatoma-derived growth factor (high-mobility group protein 1-like)	Hs.89525.0	gb:NM_004494.1	HDGF	X	2.72	0.0621
556	202466_at	topoisomerase-related function protein 4-1	Hs.225951.0	gb:NM_006999.2	TRF4	5	2.27	0.0621
557	208667_s_at	suppression of tumorigenicity 13 (colon carcinoma) (Hsp70-interacting pro	Hs.119222.1	gb:U17714.1	ST13	22	2.68	0.0621
558	204710_s_at	CGI-50 protein	Hs.279850.0	gb:NM_016003.1	LOC51623		2.51	0.0621

Fig. 19

	A	B	C	D	E	F	G	H
559	218671_s_at	ATPase inhibitor precursor	Hs.241336.0	gb:NM_016311.1	LOC5118	1	2.37	0.0622
560	217768_at	CGI-99 protein	Hs.110803.0	gb:NM_016039.1	LOC5163	14	2.76	0.0622
561	201649_at	ubiquitin-conjugating enzyme E2L 6	Hs.169895.0	gb:NM_004223.1	UBE2L6	11	2.38	0.0622
562	211858_x_at	guanine nucleotide binding protein (G protein), alpha stimulating activity p	Hs.273385.3	gb:AF088184.1		20	3.26	0.0623
563	212854_x_at	KIAA1693 protein; hypothetical protein FLJ20719	Hs.323463.0	gb:AB051480.1	KIAA169	1	2.68	0.0623
564	221452_s_at			gb:NM_030969.1	MGC122	6	2.49	0.0623
565	201103_x_at	hypothetical protein	Hs.218329.0	gb:BE299495_RC		1	2.92	0.0624
566	211960_s_at	hypothetical protein PRO2706	Hs.237955.3	gb:BG261416		3	2.16	0.0625
567	200015_s_at	neural precursor cell expressed, developmentally down-regulated 5	Hs.155595.0	gb:NM_004404.1	NEDD5	2	2.80	0.0625
568	200668_s_at	ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45)	Hs.118797.0	gb:BC003395.1		4	2.75	0.0626
569	212087_s_at	era (E. coli G-protein homolog)-like 1	Hs.3426.0	gb:AL562733_RC		17	2.13	0.0626
570	213883_s_at	glucose phosphate isomerase	Hs.180532.5	gb:AA012917_RC		1	2.27	0.0626
571	200081_s_at	ribosomal protein S6	Hs.241507.3	gb:BE741754		12	3.34	0.0626
572	211946_s_at	KIAA1096 protein	Hs.69559.2	gb:AL096857.1		1	2.39	0.0627
573	203026_at	KIAA0354 gene product	Hs.3682.0	gb:NM_014872.1	KIAA035	9	2.14	0.0627
574	201922_at	hypothetical protein	Hs.8170.0	gb:NM_014886.1	YR-29	5	2.60	0.0629
575	200845_s_at	anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium	Hs.120.0	gb:NM_004905.1	KIAA010	1	2.63	0.0629
576	200003_s_at	ribosomal protein L28	Hs.4437.0	gb:NM_000991.1	RPL28	10	3.31	0.0630
577	200940_s_at	arginine-glutamic acid dipeptide (RE) repeats	Hs.194369.0	gb:AB036737.1		1	2.50	0.0630
578	200860_s_at	KIAA1007 protein	Hs.279949.0	gb:BC000779.1		16	2.49	0.0631
579	201665_x_at	ribosomal protein S17	Hs.5174.0	gb:NM_001021.1	RPS17	15	3.47	0.0632
580	200750_s_at	RAN, member RAS oncogene family	Hs.10842.0	gb:AF054183.1		12	2.66	0.0632
581	209036_s_at	malate dehydrogenase 2, NAD (mitochondrial)	Hs.111076.0	gb:BC001917.1		7	2.79	0.0633
582	216231_s_at	beta-2-microglobulin	Hs.75415.2	gb:AW188940_RC		15	3.53	0.0633
583	208990_s_at	heterogeneous nuclear ribonucleoprotein H3 (2H9)	Hs.279681.5	gb:AF132362.1		10	2.43	0.0634
584	209112_at	cyclin-dependent kinase inhibitor 1B (p27, Kip1)	Hs.238990.0	gb:BC001971.1		12	2.55	0.0635
585	212232_at	KIAA1014 protein	Hs.6834.0	gb:AB023231.1	KIAA101	11	2.34	0.0637
586	201542_at	SAR1 protein	Hs.110796.0	gb:AY008268.1	SAR1	10	2.22	0.0637
587	207335_x_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e	Hs.85539.0	gb:NM_007100.1	ATP5I		2.40	0.0637
588	211730_s_at			gb:BC005903.1			2.75	0.0638
589	218198_at	hypothetical protein FLJ10889	Hs.171835.0	gb:NM_018180.1	FLJ10889	10	2.34	0.0638

Fig. 20

	A	B	C	D	E	F	G	H
590	202717_s_at	CDC16 (cell division cycle 16, S. cerevisiae, homolog)	Hs.1592.0	gb:NM_003903.1	CDC16	13	2.33	0.0639
591	218374_s_at	chromosome 12 open reading frame 4	Hs.296198.0	gb:NM_020374.1	C12ORF4	12	1.83	0.0639
592	217807_s_at	glioma tumor suppressor candidate region gene 2	Hs.2237.0	gb:NM_015710.1	GLTSCR2	19	3.16	0.0639
593	209046_s_at	ganglioside expression factor 2	Hs.6518.0	gb:AB030710.1	FLC3A	16	2.90	0.0641
594	213414_s_at	ribosomal protein S19	Hs.298262.1	gb:BE259729_RC		14	3.42	0.0641
595	202842_s_at	DnaJ (Hsp40) homolog, subfamily B, member 9	Hs.6790.0	gb:AL080081.1	DKFZp56	7	2.61	0.0642
596	212074_at	KIAA0810 protein	Hs.7531.2	gb:BE972774			2.46	0.0642
597	216342_x_at	Human DNA sequence from clone RP1-189G13 on chromosome 20. Conta	Hs.283838.0	gb:AL121916			3.21	0.0642
598	217748_at	CGI-45 protein	Hs.5298.0	gb:NM_015999.1	LOC5109	1	2.45	0.0643
599	208956_x_at	dUTP pyrophosphatase	Hs.82113.0	gb:U62891.1	DUT	15	2.61	0.0643
600	200909_s_at	ribosomal protein, large P2	Hs.119500.0	gb:NM_001004.1	RPLP2	11	3.35	0.0643
601	208968_s_at	hypothetical protein	Hs.4900.2	gb:BC002568.1		16	2.36	0.0644
602	217871_s_at	macrophage migration inhibitory factor (glycosylation-inhibiting factor)	Hs.73798.0	gb:NM_002415.1	MIF	22	3.06	0.0644
603	212537_x_at	ribosomal protein L17	Hs.82202.2	gb:BE733979		1	3.24	0.0644
604	211800_s_at	ubiquitin specific protease 4 (proto-oncogene)	Hs.77500.1	gb:AF017306.1	UNP	3	2.19	0.0644
605	208905_at	cytochrome c	Hs.169248.0	gb:BC005299.1		7	2.95	0.0646
606	208517_x_at	basic transcription factor 3	Hs.101025.3	gb:NM_001207.1	BTF3		3.02	0.0647
607	201375_s_at	protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform	Hs.80350.0	gb:NM_004156.1	PPP2CB	8	2.88	0.0647
608	217773_s_at	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (9kD, MLRQ)	Hs.50098.0	gb:NM_002489.1	NDUFA4	7	2.95	0.0647
609	200082_s_at	ribosomal protein S7	Hs.301547.2	gb:AI805587_RC		17	3.15	0.0648
610	220741_s_at	inorganic pyrophosphatase	Hs.5123.0	gb:NM_006903.1	SID6-306	4	2.21	0.0648
611	210125_s_at	Breakpoint cluster region protein, uterine leiomyoma, 1; barrier to autointe	Hs.268763.1	gb:AF044773.1	BCRG1	11	2.45	0.0648
612	218320_s_at	similar to mouse neuronal protein 15.6	Hs.111497.0	gb:NM_019056.1	FLJ20494		2.64	0.0648
613	205849_s_at	ubiquinol-cytochrome c reductase binding protein	Hs.131255.0	gb:NM_006294.1	UQCRB	X	2.90	0.0649
614	209944_at	clones 23667 and 23775 zinc finger protein	Hs.7137.1	gb:BC000330.1		14	2.34	0.0649
615	218082_s_at	upstream binding protein 1 (LBP-1a)	Hs.28423.0	gb:NM_014517.1	UBP1	3	2.20	0.0649
616	211047_x_at			gb:BC006337.1		19	2.67	0.0650
617	200605_s_at	protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific e	Hs.183037.0	gb:NM_002734.1	PRKAR1A	17	2.75	0.0651
618	201919_at	hypothetical protein FLJ10618	Hs.42484.0	gb:AL049246.1		3	2.58	0.0651
619	211070_x_at			gb:BC006466.1			2.82	0.0652
620	208856_x_at	ribosomal protein, large, P0	Hs.73742.1	gb:BC003655.1		2	3.55	0.0652

Fig. 21

	A	B	C	D	E	F	G	H
621	217885_at	hypothetical protein FLJ10402	Hs.7579.0	gb:NM_018085.1	FLJ10402	1	2.30	0.0652
622	201406_at	ribosomal protein L44	Hs.178391.0	gb:NM_021029.1	RPL44	X	3.31	0.0652
623	212433_x_at	ribosomal protein S2	Hs.182426.2	gb:AA630314_RC		17	3.37	0.0653
624	213011_s_at	triosephosphate isomerase 1	Hs.83848.4	gb:BF116254_RC		1	2.92	0.0653
625	217883_at	hypothetical protein	Hs.5324.0	gb:NM_015702.1	CL25022	11	2.74	0.0653
626	208929_x_at	ribosomal protein L13	Hs.180842.0	gb:BC004954.1		16	3.36	0.0653
627	204568_at	KIAA0831 protein	Hs.103000.0	gb:NM_014924.1	KIAA083	14	2.04	0.0654
628	205335_s_at	signal recognition particle 19kD	Hs.2943.0	gb:NM_003135.1	SRP19	5	2.32	0.0654
629	212270_x_at	ribosomal protein L17	Hs.82202.4	gb:BG168283		1	3.22	0.0655
630	200620_at	chromosome 1 open reading frame 8	Hs.11441.0	gb:NM_004872.1	C1ORF8	1	2.92	0.0655
631	203380_x_at	splicing factor, arginineserine-rich 5	Hs.166975.0	gb:NM_006925.1	SFRS5	14	2.55	0.0655
632	212718_at	hypothetical protein	Hs.5324.2	gb:BF797555		14	2.56	0.0655
633	221613_s_at	protein associated with PRK1	Hs.83954.3	gb:AL136598.1	DKFZp56	15	2.43	0.0655
634	202298_at	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE	Hs.74823.0	gb:NM_004541.2	NDUFA1	X	2.75	0.0655
635	202475_at	seven transmembrane domain protein	Hs.9234.0	gb:NM_006326.1	NIFIE14	19	2.57	0.0656
636	213864_s_at	nucleosome assembly protein 1-like 1	Hs.179662.3	gb:AI985751_RC		12	2.75	0.0657
637	211666_x_at			gb:L22453.1	TARBP-b	22	3.36	0.0657
638	220248_x_at	p47 protein	Hs.284261.0	gb:NM_018839.1	LOC5596	20	2.30	0.0657
639	210891_s_at	general transcription factor II, i	Hs.278589.1	gb:AF035737.1	GTF2I	7	2.73	0.0658
640	202228_s_at	stromal cell derived factor receptor 1	Hs.6354.0	gb:NM_017455.1	SDFR1	15	2.94	0.0658
641	211763_s_at			gb:BC005979.1		5	2.35	0.0658
642	202428_x_at	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A I	Hs.78888.0	gb:NM_020548.1	DBI	2	2.92	0.0659
643	208722_s_at	anaphase-promoting complex subunit 5	Hs.7101.0	gb:BC001081.1		12	2.25	0.0659
644	218592_s_at	hypothetical protein FLJ20454	Hs.26890.0	gb:NM_017829.1	FLJ20454	22	2.20	0.0659
645	208689_s_at	ribophorin II	Hs.75722.0	gb:BC003560.1		20	2.72	0.0661
646	210338_s_at	heat shock 70kD protein 8	Hs.180414.2	gb:AB034951.1	HSC54	11	3.16	0.0662
647	219762_s_at	ribosomal protein L36	Hs.300759.0	gb:NM_015414.1	RPL36	19	2.98	0.0662
648	214800_x_at	basic transcription factor 3	Hs.101025.4	gb:R83000_RC		5	2.50	0.0662
649	201317_s_at	proteasome (prosome, macropain) subunit, alpha type, 2	Hs.181309.0	gb:NM_002787.1	PSMA2	7	2.52	0.0662
650	201033_x_at	ribosomal protein, large, P0	Hs.73742.0	gb:NM_001002.1	RPLP0	12	3.55	0.0662
651	204246_s_at	dynactin 3 (p22)	Hs.15961.1	gb:NM_007234.2	DCTN3	9	2.58	0.0663

Fig. 22

	A	B	C	D	E	F	G	H
652	214281_s_at	PRO1996 protein	Hs.279864.1	gb:AA524525_RC		4	2.03	0.0664
653	213294_at	Homo sapiens clone FLC0675 PRO2870 mRNA, complete cds	Hs.306117.1	gb:AV755522		2	2.24	0.0665
654	200882_s_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	Hs.148495.0	gb:NM_002810.1	PSMD4	1	2.39	0.0666
655	201734_at	chloride channel 3	Hs.174139.0	gb:AI760629_RC		4	2.62	0.0666
656	209248_at	PTD010 protein	Hs.182470.0	gb:AL136713.1	DKFZp566K113		2.66	0.0667
657	213655_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.79474.4	gb:AA502643_RC		17	2.90	0.0667
658	202370_s_at	core-binding factor, beta subunit	Hs.179881.0	gb:NM_001755.1	CBFB	16	2.44	0.0669
659	203663_s_at	cytochrome c oxidase subunit Va	Hs.181028.0	gb:NM_004255.1	COX5A	14	2.83	0.0669
660	213465_s_at	protein phosphatase 1, regulatory subunit 7	Hs.36587.1	gb:BF718769_RC		2	2.25	0.0669
661	201441_at	cytochrome c oxidase subunit VIb	Hs.174031.0	gb:NM_001863.2	COX6B	19	2.66	0.0670
662	200610_s_at	nucleolin	Hs.79110.0	gb:NM_005381.1	NCL	2	2.82	0.0670
663	212455_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 16	Hs.12797.3	gb:N36997_RC		4	2.52	0.0672
664	211969_at	heat shock 90kD protein 1, alpha	Hs.289088.4	gb:BG420237		14	3.01	0.0672
665	201648_at	Janus kinase 1 (a protein tyrosine kinase)	Hs.50651.0	gb:AL039831_RC		1	2.59	0.0672
666	202634_at	polymerase (RNA) II (DNA directed) polypeptide K (7.0kD)	Hs.150675.0	gb:AL558030		8	1.94	0.0673
667	210950_s_at	farnesyl-diphosphate farnesyltransferase 1	Hs.48876.0	gb:BC003573.1		8	2.69	0.0673
668	209186_at	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	Hs.1526.1	gb:M23114.1	HK1	12	2.73	0.0673
669	202364_at	MAX-interacting protein 1	Hs.118630.0	gb:NM_005962.1	MXI1	10	2.75	0.0673
670	200094_s_at	eukaryotic translation elongation factor 2	Hs.75309.1	gb:AI004246_RC		19	3.13	0.0673
671	214328_s_at	heat shock 90kD protein 1, alpha	Hs.289088.3	gb:R01140_RC		14	3.23	0.0676
672	217882_at	30 kDa protein	Hs.283714.0	gb:NM_018447.1	LOC5583	3	2.41	0.0676
673	201519_at	translocase of outer mitochondrial membrane 70 (yeast) homolog A	Hs.21198.0	gb:NM_014820.1	TOMM70	3	2.54	0.0676
674	212082_s_at	myosin, light polypeptide 6, alkali, smooth muscle and non-muscle	Hs.77385.2	gb:BE734356		12	3.05	0.0678
675	219329_s_at	apoptosis related protein APR-3	Hs.9527.0	gb:NM_016085.1	3.Apr	2	2.63	0.0678
676	202649_x_at	ribosomal protein S19	Hs.298262.0	gb:NM_001022.1	RPS19	19	3.33	0.0678
677	200834_s_at	ribosomal protein S21	Hs.1948.0	gb:NM_001024.1	RPS21	20	3.27	0.0679
678	212416_at	secretory carrier membrane protein 1	Hs.31218.1	gb:AV745949		5	2.36	0.0679
679	211720_x_at			gb:BC005863.1		12	3.54	0.0680
680	200948_at	myeloid leukemia factor 2	Hs.79026.0	gb:NM_005439.1	MLF2	12	2.51	0.0680
681	201096_s_at	ADP-ribosylation factor 4	Hs.75290.0	gb:AL537042		3	2.51	0.0681
682	202753_at	KIAA0107 gene product	Hs.23488.0	gb:NM_014814.1	KIAA010	3	2.56	0.0681

Fig. 23

	A	B	C	D	E	F	G	H
683	210638_s_at	F-box only protein 9	Hs.11050.1	gb:AF176704.1		6	2.55	0.0682
684	202683_s_at	RNA (guanine-7-) methyltransferase	Hs.8086.0	gb:NM_003799.1	RNMT	18	2.09	0.0682
685	200655_s_at	calmodulin 1 (phosphorylase kinase, delta)	Hs.177656.0	gb:NM_006888.1	CALM1	14	2.92	0.0683
686	218203_at	Alg5, S. cerevisiae, homolog of	Hs.227933.0	gb:NM_013338.2	ALG5	13	2.41	0.0684
687	201738_at	translation factor sui1 homolog	Hs.21756.0	gb:NM_005875.1	GC20	3	2.66	0.0684
688	205690_s_at	maternal G10 transcript	Hs.330310.0	gb:NM_003910.1	G10	7	2.10	0.0684
689	201577_at	non-metastatic cells 1, protein (NM23A) expressed in	Hs.118638.0	gb:NM_000269.1	NME1	17	2.50	0.0686
690	211747_s_at			gb:BC005938.1		7	2.19	0.0688
691	218563_at	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9)	Hs.198269.0	gb:NM_004542.1	NDUFA3	19	2.57	0.0688
692	213229_at	helicase-moi	Hs.87889.1	gb:BF590131_RC		14	2.40	0.0688
693	215691_x_at	HSPCO34 protein	Hs.46967.2	gb:AV702994		1	2.20	0.0692
694	215158_s_at	death effector domain-containing	Hs.169681.2	gb:AK022531.1			2.15	0.0692
695	212578_x_at	ribosomal protein S17	Hs.5174.2	gb:BF026595		4	3.34	0.0692
696	217918_at	HSPC162 protein	Hs.100002.0	gb:NM_014183.1	HSPC162	20	2.85	0.0693
697	214882_s_at	splicing factor, arginineserine-rich 2	Hs.73965.1	gb:BG254869		17	2.48	0.0693
698	202907_s_at	Nijmegen breakage syndrome 1 (nibrin)	Hs.25812.0	gb:NM_002485.2	NBS1	8	2.23	0.0694
699	202277_at	serine palmitoyltransferase, long chain base subunit 1	Hs.90458.0	gb:AL568804_RC		9	2.39	0.0694
700	202136_at	adenovirus 5 E1A binding protein	Hs.301449.0	gb:BE250417_RC		10	2.58	0.0695
701	202001_s_at	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)	Hs.274416.0	gb:NM_002490.1	NDUFA6	22	2.51	0.0695
702	212144_at	KIAA0668 protein	Hs.5898.1	gb:AL021707		22	2.29	0.0696
703	202276_at	Deleted in split-handsplit-foot 1 region	Hs.85215.0	gb:NM_006304.1	DSS1	7	2.37	0.0696
704	219206_x_at	CGI-119 protein	Hs.283670.0	gb:NM_016056.1	LOC5164	12	2.49	0.0698
705	213187_x_at	alpha-2-HS-glycoprotein	Hs.324746.2	gb:BG538564		19	3.31	0.0700
706	221475_s_at	ribosomal protein L15	Hs.74267.0	gb:NM_002948.1	RPL15	3	3.11	0.0700
707	200850_s_at	S-adenosylhomocysteine hydrolase-like 1	Hs.4113.0	gb:NM_006621.1	AHCYL1	1	2.81	0.0700
708	201003_x_at	ubiquitin-conjugating enzyme E2 variant 1	Hs.75875.0	gb:NM_003349.2	UBE2V1	20	2.40	0.0702
709	200777_s_at	KIAA0005 gene product	Hs.155291.0	gb:NM_014670.1	KIAA000	2	2.47	0.0703
710	208818_s_at	catechol-O-methyltransferase	Hs.240013.0	gb:BC000419.1		22	2.61	0.0703
711	211115_x_at	survival of motor neuron protein interacting protein 1	Hs.102456.3	gb:AB037703.1	SIP1-delta	14	1.94	0.0703
712	200807_s_at	heat shock 60kD protein 1 (chaperonin)	Hs.79037.0	gb:NM_002156.1	HSPD1	2	3.05	0.0703
713	206158_s_at	zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)	Hs.2110.0	gb:NM_003418.1	ZNF9	3	2.73	0.0704

Fig. 24

	A	B	C	D	E	F	G	H
714	202811_at	associated molecule with the SH3 domain of STAM	Hs.12479.0	gb:NM_006463.1	AMSH	2	2.15	0.0704
715	219939_s_at	NRAS-related gene	Hs.69855.0	gb:NM_007158.1	D1S155E	1	2.78	0.0705
716	208726_s_at	eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD)	Hs.12163.0	gb:BC000461.1		20	2.67	0.0705
717	217753_s_at	ribosomal protein S26	Hs.299465.0	gb:NM_001029.1	RPS26	12	3.05	0.0705
718	203545_at	hypothetical protein MGC2840 similar to a putative glucosyltransferase	Hs.155356.0	gb:NM_024079.1	MGC284(	11	2.24	0.0706
719	200087_s_at	Homo sapiens cDNA: FLJ21323 fis, clone COL02374	Hs.75914.0	gb:AK024976.1		12	2.81	0.0707
720	202941_at	NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD)	Hs.51299.0	gb:NM_021074.1	NDUFV2	18	2.51	0.0707
721	210283_x_at	polyadenylate binding protein-interacting protein 1	Hs.109643.1	gb:BC005295.1		17	2.30	0.0708
722	209481_at	KIAA0096 protein	Hs.79025.0	gb:AF226044.1	HSNFRK	3	2.41	0.0708
723	214042_s_at	Homo sapiens mRNA; cDNA DKFZp586E0524 (from clone DKFZp586E0	Hs.326249.0	gb:AW071997_RC		X	2.57	0.0709
724	211452_x_at	leucine rich repeat (in FLII) interacting protein 1	Hs.326159.1	gb:AF130054.1_RC		2	2.89	0.0710
725	204031_s_at	poly(rC)-binding protein 2	Hs.63525.0	gb:NM_005016.1	PCBP2	12	2.98	0.0710
726	200090_at	farnesyltransferase, CAAX box, alpha	Hs.138381.1	gb:BG168896		13	2.66	0.0710
727	212454_x_at	heterogeneous nuclear ribonucleoprotein D-like	Hs.170311.2	gb:AI762552_RC		4	2.66	0.0712
728	221543_s_at	chromosome 8 open reading frame 2	Hs.125849.0	gb:AL442077.1	DKFZp66	8	2.08	0.0712
729	200858_s_at	ribosomal protein S8	Hs.151604.0	gb:NM_001012.1	RPS8	5	3.09	0.0712
730	200088_x_at	ribosomal protein L12	Hs.182979.1	gb:AK026491.1		2	3.37	0.0712
731	202141_s_at	COP9 homolog	Hs.75193.0	gb:BC003090.1		2	2.47	0.0713
732	218027_at	HSPC145 protein	Hs.18349.0	gb:NM_014175.1	HSPC145	8	2.36	0.0713
733	221570_s_at	HSPC133 protein	Hs.273063.1	gb:AF201938.1	DC3	2	2.28	0.0713
734	212595_s_at	DAZ associated protein 2	Hs.75416.1	gb:AL534321		12	2.55	0.0713
735	218171_at	suppressor of K+ transport defect 1	Hs.126550.0	gb:AF195514.1	VPS42	18	2.43	0.0714
736	212226_s_at	Homo sapiens phosphatidic acid phosphatase type 2B (PPAP2B), mRNA	Hs.173717.0	gb:AA628586_RC		1	2.84	0.0714
737	210912_x_at	glutathione S-transferase M4	Hs.301961.1	gb:M99422.1	GST	1	2.17	0.0715
738	200618_at	LIM and SH3 protein 1	Hs.75080.0	gb:NM_006148.1	LASP1	17	2.59	0.0716
739	206792_x_at	phosphodiesterase 4C, cAMP-specific (dunce (Drosophila)-homolog phosp	Hs.189.0	gb:NM_000923.1	PDE4C	19	2.76	0.0716
740	201210_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 3	Hs.147916.0	gb:NM_001356.2	DDX3	X	2.61	0.0717
741	209630_s_at	G protein-coupled receptor 37 (endothelin receptor type B-like)	Hs.27747.0	gb:AL043967_RC		9	1.98	0.0717
742	202594_at	leptin receptor overlapping transcript-like 1	Hs.11000.0	gb:NM_015344.1	MY047	8	2.35	0.0718
743	201947_s_at	chaperonin containing TCP1, subunit 2 (beta)	Hs.6456.0	gb:NM_006431.1	CCT2	12	2.58	0.0720
744	212191_x_at	ribosomal protein L13	Hs.180842.3	gb:AW574664_RC		9	3.30	0.0720

Fig. 25

	A	B	C	D	E	F	G	H
745	203320_at	lymphocyte adaptor protein	Hs.13131.0	gb:NM_005475.1	LNK	12	2.12	0.0721
746	200899_s_at	meningioma expressed antigen 5 (hyaluronidase)	Hs.5734.0	gb:NM_012215.1	MGEA5	10	2.33	0.0721
747	201268_at	non-metastatic cells 2, protein (NM23B) expressed in	Hs.275163.0	gb:NM_002512.1	NME2	17	2.75	0.0721
748	217747_s_at	ribosomal protein S9	Hs.180920.0	gb:NM_001013.1	RPS9	19	3.10	0.0722
749	210406_s_at	RAB6A, member RAS oncogene family	Hs.5636.1	gb:AL136727.1	DKFZp56	2	2.50	0.0723
750	210211_s_at	heat shock 90kD protein 1, alpha	Hs.289088.0	gb:AF028832.1		14	2.99	0.0724
751	201226_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI)	Hs.198273.0	gb:NM_005004.1	NDUFB8	10	2.68	0.0724
752	201773_at	activity-dependent neuroprotective protein	Hs.3657.0	gb:NM_015339.1	ADNP	20	2.33	0.0725
753	208093_s_at			gb:NM_030808.1	NUDEL	17	2.32	0.0725
754	216032_s_at	hypothetical 43.2 Kd protein	Hs.169992.1	gb:AF091085.1		20	2.53	0.0725
755	218659_at	KIAA1685 protein	Hs.13801.0	gb:NM_018263.1	FLJ10898	2	2.38	0.0725
756	201586_s_at	splicing factor prolineglutamine rich (polypyrimidine tract-binding protein-	Hs.180610.0	gb:NM_005066.1	SFPQ	1	2.41	0.0726
757	211988_at	SWISNF related, matrix associated, actin dependent regulator of chromatin	Hs.332848.0	gb:BG289800		17	2.31	0.0727
758	205133_s_at	heat shock 10kD protein 1 (chaperonin 10)	Hs.1197.0	gb:NM_002157.1	HSPE1	2	2.35	0.0728
759	218946_at	HIRIP5 protein; HIRA-interacting protein 5	Hs.17368.0	gb:NM_015700.1	HIRIP5		2.02	0.0728
760	201306_s_at	acidic protein rich in leucines	Hs.84264.0	gb:NM_006401.1	SSP29	9	2.95	0.0728
761	221493_at	TSPY-like	Hs.278479.0	gb:AL136629.1	DKFZp56	6	2.57	0.0728
762	201200_at	cellular repressor of E1A-stimulated genes	Hs.5710.0	gb:NM_003851.1	CREG	1	2.59	0.0729
763	208799_at	proteasome (prosome, macropain) subunit, beta type, 5	Hs.78596.0	gb:BC004146.1		14	2.52	0.0730
764	213911_s_at	H2A histone family, member Z	Hs.119192.1	gb:BF718636_RC		12	2.68	0.0730
765	201054_at	heterogeneous nuclear ribonucleoprotein A0	Hs.77492.0	gb:BE966599_RC		5	2.48	0.0730
766	203460_s_at	presenilin 1 (Alzheimer disease 3)	Hs.3260.0	gb:NM_007318.1	PSEN1	14	2.21	0.0730
767	200052_s_at	interleukin enhancer binding factor 2, 45kD	Hs.75117.0	gb:NM_004515.1	ILF2	14	2.33	0.0735
768	210835_s_at	C-terminal binding protein 2	Hs.171391.1	gb:AF222711.1		10	2.64	0.0735
769	200996_at	ARP3 (actin-related protein 3, yeast) homolog	Hs.5321.0	gb:NM_005721.2	ACTR3	2	2.53	0.0735
770	217803_at	Golgi protein	Hs.18271.0	gb:NM_022130.1	GPP34	5	2.50	0.0736
771	209066_x_at	ubiquinol-cytochrome c reductase binding protein	Hs.131255.1	gb:M26700.1	UQBP	8	2.81	0.0736
772	215823_x_at	poly(A)-binding protein, cytoplasmic, pseudogene 3	Hs.283767.0	gb:U64661		X	3.02	0.0737
773	212245_at	Human transposon-like element mRNA	Hs.84775.0	gb:AL567779_RC		2	2.28	0.0738
774	213102_at	hypothetical protein EUROIMAGE1875335	Hs.10927.1	gb:Z78330		2	2.52	0.0739
775	200072_s_at	heterogeneous nuclear ribonucleoprotein M	Hs.79024.1	gb:AF061832.1		NA	2.52	0.0739

Fig. 26

	A	B	C	D	E	F	G	H
776	208666_s_at	suppression of tumorigenicity 13 (colon carcinoma) (Hsp70-interacting pro	Hs.119222.1	gb:BE866412		22	2.32	0.0741
777	200809_x_at	ribosomal protein L12	Hs.182979.0	gb:NM_000976.1	RPL12	9	3.35	0.0743
778	201023_at	TATA box binding protein (TBP)-associated factor, RNA polymerase II, F	Hs.155188.0	gb:NM_005642.1	TAF2F	5	2.44	0.0744
779	211972_x_at	ribosomal protein, large, P0	Hs.73742.3	gb:AI953822_RC		5	3.42	0.0744
780	208647_at	farnesyl-diphosphate farnesyltransferase 1	Hs.48876.1	gb:AA872727_RC		8	2.62	0.0745
781	201470_at	glutathione-S-transferase like; glutathione transferase omega	Hs.11465.0	gb:NM_004832.1	GSTTLp2	10	2.74	0.0745
782	200799_at	heat shock 70kD protein 1A	Hs.8997.0	gb:NM_005345.3	HSPA1A	6	3.07	0.0746
783	208679_s_at	actin related protein 23 complex, subunit 2 (34 kD)	Hs.83583.1	gb:AF279893.1		2	2.80	0.0746
784	217731_s_at	integral membrane protein 2B	Hs.239625.0	gb:NM_021999.1	ITM2B	13	2.79	0.0746
785	209069_s_at	H3 histone, family 3A	Hs.181307.1	gb:BC001124.1		17	2.92	0.0746
786	200662_s_at	translocase of outer mitochondrial membrane 20 (yeast) homolog	Hs.75187.0	gb:NM_014765.1	KIAA001	X	2.68	0.0747
787	200711_s_at	transcription elongation factor B (SIII), polypeptide 1-like	Hs.171626.0	gb:NM_003197.2	TCEB1L	5	2.75	0.0747
788	200975_at	palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile	Hs.3873.0	gb:NM_000310.1	PPT1		2.64	0.0747
789	203137_at	Wilms tumour 1-associating protein	Hs.119.0	gb:NM_004906.1	KIAA010	6	2.24	0.0749
790	208152_s_at			gb:NM_004728.1	DDX21	10	2.38	0.0750
791	211938_at	hypothetical protein PRO1843	Hs.283330.1	gb:BF247371		12	2.59	0.0751
792	215096_s_at	esterase Dformylglutathione hydrolase	Hs.82193.1	gb:AU145746_RC		13	2.53	0.0752
793	201290_at	signal peptidase complex (18kD)	Hs.9534.0	gb:NM_014300.1	SPC18	15	2.60	0.0753
794	221532_s_at	recombination protein REC14	Hs.296242.0	gb:AF309553.1		15	2.29	0.0753
795	210024_s_at	ubiquitin-conjugating enzyme E2E 3 (homologous to yeast UBC45)	Hs.4890.0	gb:AB017644.1	UbcH9	X	2.48	0.0753
796	212888_at	Homo sapiens clone 23938 mRNA sequence	Hs.325625.0	gb:BG109746		14	2.23	0.0754
797	217869_at	steroid dehydrogenase homolog	Hs.279617.0	gb:NM_016142.1	LOC5114	11	2.59	0.0756
798	200989_at	hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcripti	Hs.197540.0	gb:NM_001530.1	HIF1A	14	2.72	0.0756
799	208986_at	transcription factor 12 (HTF4, helix-loop-helix transcription factors 4)	Hs.21704.0	gb:AL559478		15	2.37	0.0756
800	200972_at	tetraspan 3	Hs.100090.0	gb:BC000704.1			2.79	0.0757
801	216438_s_at	sterol-C5-desaturase (fungal ERG3, delta-5-desaturase)-like	Hs.288031.2	gb:AL133228			3.13	0.0757
802	201216_at	endoplasmic reticulum lumenal protein	Hs.75841.0	gb:NM_006817.2	ERP28	12	2.63	0.0757
803	200673_at	lysosomal-associated protein transmembrane 4 alpha	Hs.111894.0	gb:NM_014713.2	MBNT	2	2.91	0.0757
804	218117_at	ring-box 1	Hs.279919.0	gb:NM_014248.1	RBX1	22	2.36	0.0759
805	202675_at	succinate dehydrogenase complex, subunit B, iron sulfur (Ip)	Hs.64.0	gb:NM_003000.1	SDHB	1	2.13	0.0759
806	211025_x_at			gb:BC006229.1			2.69	0.0760

Fig. 27

	A	B	C	D	E	F	G	H
807	215380_s_at	hypothetical protein MGC3077	Hs.288649.2	gb:AK021779.1		7	2.36	0.0762
808	210825_s_at	erythrocyte membrane protein band 7.2 (stomatin)	Hs.160483.1	gb:AF130103.1_RC		9	3.07	0.0764
809	208639_x_at	protein disulfide isomerase-related protein	Hs.182429.1	gb:BC001312.1		2	2.72	0.0764
810	210111_s_at	Homo sapiens PNAS-138 mRNA, complete cds	Hs.326790.0	gb:AF277175.1		7	2.38	0.0765
811	200985_s_at	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5	Hs.119663.0	gb:NM_000611.1	CD59	11	2.74	0.0765
812	207668_x_at	protein disulfide isomerase-related protein	Hs.182429.0	gb:NM_005742.1	P5	2	2.80	0.0765
813	201612_at	aldehyde dehydrogenase 9 family, member A1	Hs.2533.0	gb:NM_000696.1	ALDH9	1	2.57	0.0765
814	215313_x_at	major histocompatibility complex, class I, A	Hs.181244.4	gb:AA573862_RC		6	3.38	0.0765
815	201013_s_at	multifunctional polypeptide similar to SAICAR synthetase and AIR carboxy	Hs.117950.0	gb:AA902652_RC		4	2.31	0.0766
816	206874_s_at	Ste20-related serinethreonine kinase	Hs.105751.0	gb:AL138761		10	2.23	0.0766
817	200075_s_at			gb:BC006249.1		1	2.51	0.0766
818	211623_s_at			gb:M30448.1	CSNK2B		2.58	0.0766
819	214041_x_at	ribosomal protein L37a	Hs.5566.3	gb:BE857772_RC			2.20	0.0767
820	200761_s_at	vitamin A responsive; cytoskeleton related	Hs.92384.0	gb:NM_006407.2	JWA	3	2.90	0.0767
821	217870_s_at	UMP-CMP kinase	Hs.11463.0	gb:NM_016308.1	LOC5172	1	2.51	0.0768
822	217800_s_at	hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5	Hs.9788.0	gb:NM_030571.1	MGC109	5	2.65	0.0768
823	200764_s_at	catenin (cadherin-associated protein), alpha 1 (102kD)	Hs.178452.0	gb:AI826881_RC		5	2.28	0.0769
824	201916_s_at	SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like	Hs.31575.0	gb:NM_007214.1	SEC63L	6	2.36	0.0771
825	201238_s_at	capping protein (actin filament) muscle Z-line, alpha 2	Hs.75546.0	gb:BC005338.1		7	2.41	0.0772
826	212205_at	purine-rich element binding protein B	Hs.301005.1	gb:AA534860_RC		7	2.60	0.0773
827	211971_s_at	leucine-rich protein mRNA	Hs.182490.0	gb:AI653608_RC			2.38	0.0773
828	208671_at	KIAA1253 protein	Hs.146668.0	gb:AF164794.1		6	2.72	0.0773
829	201619_at	peroxiredoxin 3	Hs.75454.0	gb:NM_006793.1	PRDX3	10	2.57	0.0773
830	200886_s_at	phosphoglycerate mutase 1 (brain)	Hs.181013.0	gb:NM_002629.1	PGAM1	10	2.90	0.0773
831	209063_x_at	polyadenylate binding protein-interacting protein 1	Hs.109643.2	gb:BF248165		17	2.28	0.0774
832	212295_s_at	Homo sapiens mRNA; cDNA DKFZp564D016 (from clone DKFZp564D0	Hs.14846.0	gb:AA148507_RC		13	2.49	0.0774
833	207941_s_at	splicing factor (CC1.3)	Hs.145696.0	gb:NM_004902.1	CC1.3	20	2.57	0.0774
834	215001_s_at	glutamate-ammonia ligase (glutamine synthase)	Hs.170171.1	gb:AL161952.1	DKFZp43	1	3.09	0.0774
835	208703_s_at	amyloid beta (A4) precursor-like protein 2	Hs.279518.1	gb:BG427393		11	2.56	0.0774
836	221702_s_at			gb:AF353992.1	BLP2	15	2.52	0.0775
837	214359_s_at	heat shock 90kD protein 1, beta	Hs.74335.1	gb:AI218219_RC		6	2.73	0.0775

Fig. 28

	A	B	C	D	E	F	G	H
838	216304_x_at	YME1 (S.cerevisiae)-like 1	Hs.206521.1	gb:AJ295618		10	2.35	0.0775
839	200824_at	glutathione S-transferase pi	Hs.226795.0	gb:NM_000852.2	GSTP1	11	2.64	0.0776
840	201877_s_at	protein phosphatase 2, regulatory subunit B (B56), gamma isoform	Hs.171734.0	gb:NM_002719.1	PPP2R5C	14	2.36	0.0776
841	213046_at	poly(A)-binding protein, nuclear 1	Hs.117176.1	gb:AI130920_RC		14	2.33	0.0777
842	211749_s_at			gb:BC005941.1		1	2.49	0.0777
843	211026_s_at			gb:BC006230.1			2.64	0.0778
844	212426_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.74405.1	gb:BF033313		2	2.87	0.0778
845	200887_s_at	signal transducer and activator of transcription 1, 91kD	Hs.21486.0	gb:NM_007315.1	STAT1	2	2.44	0.0781
846	209563_x_at	calmodulin 2 (phosphorylase kinase, delta)	Hs.182278.1	gb:BC000454.1		14	2.94	0.0783
847	211727_s_at			gb:BC005895.1		17	2.16	0.0784
848	215111_s_at	transforming growth factor beta-stimulated protein TSC-22	Hs.114360.1	gb:AK027071.1		13	3.03	0.0785
849	200903_s_at	S-adenosylhomocysteine hydrolase	Hs.172673.0	gb:NM_000687.1	AHCY	20	2.31	0.0786
850	203113_s_at	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchang	Hs.223241.0	gb:NM_001960.1	EEF1D	8	2.84	0.0786
851	211941_s_at	prostatic binding protein	Hs.80423.1	gb:BE969671		12	3.11	0.0787
852	203455_s_at	spermidinespermine N1-acetyltransferase	Hs.28491.0	gb:NM_002970.1	SAT		2.73	0.0789
853	212674_s_at	Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990	Hs.281616.0	gb:AK002076.1		3	2.20	0.0789
854	200036_s_at	ribosomal protein L10a	Hs.252574.0	gb:NM_007104.2	RPL10A	6	3.04	0.0791
855	200725_x_at	ribosomal protein L10	Hs.29797.0	gb:NM_006013.1	RPL10	X	3.38	0.0792
856	213857_s_at	CD47 antigen (Rh-related antigen, integrin-associated signal transducer)	Hs.82685.1	gb:BG230614_RC		3	2.49	0.0792
857	222231_s_at	hypothetical protein PRO1855	Hs.283558.4	gb:AK025328.1		17	2.51	0.0793
858	202026_at	succinate dehydrogenase complex, subunit D, integral membrane protein	Hs.168289.0	gb:NM_003002.1	SDHD	11	2.39	0.0794
859	201351_s_at	YME1 (S.cerevisiae)-like 1	Hs.206521.0	gb:AF070656.1		10	2.36	0.0795
860	212168_at	putative brain nuclearly-targeted protein	Hs.180895.0	gb:AL514547_RC		20	2.06	0.0795
861	202097_at	nucleoporin 153kD	Hs.211608.0	gb:NM_005124.1	NUP153	6	2.19	0.0795
862	200743_s_at	ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disea	Hs.20478.0	gb:NM_000391.2	CLN2	11	2.63	0.0795
863	217845_x_at	DKFZP564K247 protein	Hs.7917.0	gb:NM_014056.1	DKFZP56	12	2.72	0.0796
864	212870_at	son of sevenless (Drosophila) homolog 1	Hs.326392.0	gb:AI628605_RC		14	2.19	0.0796
865	200888_s_at	ribosomal protein L23	Hs.234518.0	gb:NM_000978.1	RPL23	17	3.18	0.0796
866	202614_at	chromosome 4 open reading frame 1	Hs.270956.0	gb:NM_006345.2	C4ORF1	4	2.11	0.0797
867	220939_s_at	hypothetical protein FLJ20283	Hs.44033.0	gb:NM_017743.1	FLJ20283	15	2.24	0.0798
868	212229_s_at	F-box only protein 21	Hs.184227.1	gb:AK001699.1		12	2.54	0.0798

Fig. 29

	A	B	C	D	E	F	G	H
869	214363_s_at	matrin 3	Hs.78825.2	gb:AA129420_RC		4	2.89	0.0798
870	202646_s_at	neuroblastoma RAS viral (v-ras) oncogene homolog	Hs.260523.0	gb:AA167775_RC		1	2.69	0.0799
871	212616_at	KIAA0308 protein	Hs.10351.0	gb:BF668950		16	2.27	0.0800
872	212652_s_at	sorting nexin 4	Hs.267812.1	gb:AA524345_RC		3	2.26	0.0801
873	212377_s_at	Notch (Drosophila) homolog 2	Hs.8121.3	gb:AU158495_RC		1	2.30	0.0802
874	200651_at	guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1	Hs.5662.0	gb:NM_006098.1	GNB2L1	5	3.23	0.0802
875	200936_at	ribosomal protein L8	Hs.178551.0	gb:NM_000973.1	RPL8	8	3.29	0.0804
876	208743_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.279920.1	gb:BC001359.1		20	2.26	0.0805
877	200883_at	ubiquinol-cytochrome c reductase core protein II	Hs.173554.0	gb:NM_003366.1	UQCRC2	16	2.39	0.0808
878	200984_s_at	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5,	Hs.119663.0	gb:X16447.1		11	2.77	0.0808
879	205644_s_at	small nuclear ribonucleoprotein polypeptide G	Hs.77496.0	gb:NM_003096.1	SNRPG	NA	2.40	0.0808
880	202172_at	zinc finger protein 161	Hs.6557.0	gb:BG035116		17	2.27	0.0809
881	202678_at	general transcription factor IIA, 2 (12kD subunit)	Hs.76362.0	gb:NM_004492.1	GTF2A2	15	2.37	0.0810
882	211783_s_at			gb:BC006177.1			2.20	0.0810
883	200024_at	ribosomal protein S5	Hs.76194.0	gb:NM_001009.1	RPS5	5	3.17	0.0811
884	201999_s_at	t-complex-associated-testis-expressed 1-like 1	Hs.266940.0	gb:NM_006519.1	TCTEL1	6	2.59	0.0812
885	201921_at	guanine nucleotide binding protein 10	Hs.79126.0	gb:NM_004125.1	GNG10	9	2.42	0.0813
886	201772_at	antizyme inhibitor	Hs.223014.0	gb:NM_015878.1	LOC5158	8	2.18	0.0814
887	207761_s_at	DKFZP586A0522 protein	Hs.288771.0	gb:NM_014033.1	DKFZP58	12	2.87	0.0815
888	213327_s_at	ubiquitin specific protease 12	Hs.42400.0	gb:AI820101_RC		5	1.89	0.0816
889	201347_x_at	glyoxylate reductasehydroxypyruvate reductase	Hs.155742.0	gb:NM_012203.1	GRHPR	9	2.41	0.0817
890	217168_s_at	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-l	Hs.146393.1	gb:AF217990.1		16	2.73	0.0818
891	200638_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.75103.0	gb:BC003623.1		8	2.63	0.0819
892	212034_s_at	KIAA1067 protein	Hs.325530.0	gb:AB028990.1	KIAA106	17	2.28	0.0820
893	201163_s_at	insulin-like growth factor binding protein 7	Hs.119206.0	gb:NM_001553.1	IGFBP7	4	2.83	0.0820
894	215884_s_at	ubiquilin 2	Hs.4552.1	gb:AK001029.1		X	2.16	0.0821
895	211787_s_at			gb:BC006210.1		X	2.97	0.0821
896	218605_at	hypothetical protein FLJ23182	Hs.7395.0	gb:NM_022366.1	FLJ23182	1	2.01	0.0822
897	208847_s_at	alcohol dehydrogenase 5 (class III), chi polypeptide	Hs.78989.0	gb:M29872.1	ADH5	4	2.42	0.0824
898	201587_s_at	interleukin-1 receptor-associated kinase 1	Hs.182018.0	gb:NM_001569.2	IRAK1	X	2.31	0.0824
899	213005_s_at	KIAA0172 protein	Hs.77546.0	gb:D79994.1	KIAA017	9	2.45	0.0827

Fig. 30

	A	B	C	D	E	F	G	H
900	209067_s_at	heterogeneous nuclear ribonucleoprotein D-like	Hs.170311.1	gb:D89092.1	hnRNP JK	4	2.33	0.0828
901	200944_s_at	high-mobility group (nonhistone chromosomal) protein 14	Hs.251064.0	gb:NM_004965.1	HMG14	21	2.70	0.0829
902	212050_at	Homo sapiens cDNA: FLJ23260 fis, clone COL05804, highly similar to HS	Hs.13996.0	gb:AK026913.1		17	2.29	0.0830
903	208812_x_at	major histocompatibility complex, class I, C	Hs.277477.0	gb:BC004489.1		6	3.36	0.0830
904	213969_x_at	ribosomal protein L29	Hs.183698.2	gb:BF683426		3	3.21	0.0831
905	221505_at	hypothetical protein MGC5350	Hs.71331.0	gb:AW612574_RC		1	2.21	0.0834
906	201477_s_at	ribonucleotide reductase M1 polypeptide	Hs.2934.0	gb:NM_001033.1	RRM1	11	2.19	0.0835
907	221844_x_at	hypothetical protein FLJ22059	Hs.13323.1	gb:AV756161		4	2.52	0.0836
908	202231_at	dendritic cell protein	Hs.69469.0	gb:NM_006360.1	GA17	11	2.44	0.0837
909	208405_s_at	CD164 antigen, sialomucin	Hs.43910.0	gb:NM_006016.1	CD164	6	2.78	0.0839
910	208731_at	RAB2, member RAS oncogene family	Hs.78305.0	gb:AU158062_RC		8	2.53	0.0839
911	212725_s_at	peptidylprolyl isomerase B (cyclophilin B)	Hs.699.2	gb:N37081_RC		22	2.45	0.0840
912	202362_at	RAP1A, member of RAS oncogene family	Hs.865.0	gb:NM_002884.1	RAP1A	1	2.54	0.0841
913	201779_s_at	ring finger protein 13	Hs.6900.0	gb:AF070558.1		3	2.55	0.0841
914	208669_s_at	CREBBPEP300 inhibitory protein 1	Hs.75847.0	gb:AF109873.1		15	2.53	0.0842
915	208073_x_at	tetratricopeptide repeat domain 3	Hs.118174.0	gb:NM_003316.1	TTC3	21	2.85	0.0844
916	218093_s_at	hypothetical protein FLJ20093	Hs.172572.0	gb:NM_017664.1	FLJ20093	13	2.19	0.0845
917	216526_x_at	major histocompatibility complex, class I, C	Hs.277477.2	gb:AK024836.1		6	3.40	0.0846
918	207585_s_at	ribosomal protein L36a	Hs.118857.0	gb:NM_001001.1	RPL36A	14	2.78	0.0848
919	202377_at	leptin receptor gene-related protein	Hs.23581.0	gb:AW026535_RC		1	2.56	0.0848
920	208736_at	actin related protein 23 complex, subunit 3 (21 kD)	Hs.6895.0	gb:AF004561.1		12	2.40	0.0850
921	213564_x_at	lactate dehydrogenase B	Hs.234489.1	gb:BE042354_RC		X	3.10	0.0850
922	208374_s_at	capping protein (actin filament) muscle Z-line, alpha 1	Hs.184270.0	gb:NM_006135.1	CAPZA1	1	2.47	0.0850
923	201443_s_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump) membrane sec	Hs.183434.0	gb:AF248966.1		X	2.67	0.0851
924	200666_s_at	DnaJ (Hsp40) homolog, subfamily B, member 1	Hs.82646.0	gb:NM_006145.1	HSPF1	NA	2.50	0.0851
925	201980_s_at	Homo sapiens Ras suppressor protein 1 (RSU1), mRNA	Hs.75551.0	gb:NM_012425.2	RSU1	10	2.08	0.0852
926	208692_at	ribosomal protein S3	Hs.252259.0	gb:U14990.1	rpS3	11	3.27	0.0853
927	201300_s_at	prion protein (p27-30) (Creutzfeld-Jakob disease, Gerstmann-Strausler-Sch	Hs.74621.0	gb:NM_000311.1	PRNP	20	2.74	0.0853
928	202011_at	tight junction protein 1 (zona occludens 1)	Hs.74614.0	gb:NM_003257.1	TJP1	15	2.53	0.0853
929	208683_at	calpain 2, (mII) large subunit	Hs.76288.0	gb:M23254.1	CANP	1	2.86	0.0854
930	201857_at	M-phase phosphoprotein homolog	Hs.173518.0	gb:NM_016107.1	LOC5166	5	2.50	0.0855

Fig. 31

	A	B	C	D	E	F	G	H
931	205361_s_at	prefoldin 4	Hs.91161.0	gb:AI718295_RC		20	1.95	0.0856
932	203132_at	retinoblastoma 1 (including osteosarcoma)	Hs.75770.0	gb:NM_000321.1	RB1	13	1.92	0.0859
933	214665_s_at	calcium binding protein P22	Hs.85301.3	gb:AK000095.1		15	2.49	0.0859
934	201129_at	splicing factor, arginineserine-rich 7 (35kD)	Hs.184167.0	gb:NM_006276.2	SFRS7	2	2.25	0.0861
935	200978_at	malate dehydrogenase 1, NAD (soluble)	Hs.75375.0	gb:NM_005917.1	MDH1	2	2.91	0.0862
936	203594_at	RNA 3-terminal phosphate cyclase	Hs.27076.0	gb:NM_003729.1	RPC	1	1.94	0.0862
937	200853_at	H2A histone family, member Z	Hs.119192.0	gb:NM_002106.1	H2AFZ	4	2.50	0.0863
938	202794_at	inositol polyphosphate-1-phosphatase	Hs.32309.0	gb:NM_002194.2	INPP1	2	2.45	0.0863
939	208767_s_at	Homo sapiens mRNA; cDNA DKFZp586E1124 (from clone DKFZp586E1	Hs.296398.0	gb:AW149681_RC		4	2.57	0.0865
940	207791_s_at	RAB1, member RAS oncogene family	Hs.3642.0	gb:NM_004161.1	RAB1	2	2.15	0.0865
941	203538_at	calcium modulating ligand	Hs.13572.0	gb:NM_001745.1	CAMLG	5	2.29	0.0867
942	209092_s_at	CGI-150 protein	Hs.279061.0	gb:AF061730.1		17	2.17	0.0867
943	203935_at	activin A receptor, type I	Hs.150402.0	gb:NM_001105.2	ACVR1	2	2.10	0.0868
944	218092_s_at	HIV-1 Rev binding protein	Hs.171545.0	gb:NM_004504.2	HRB		2.06	0.0869
945	202690_s_at	small nuclear ribonucleoprotein D1 polypeptide (16kD)	Hs.86948.0	gb:BC001721.1		18	2.06	0.0870
946	200742_s_at	ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disea	Hs.20478.0	gb:BG231932_RC		11	2.47	0.0870
947	211600_at			gb:U20489.1	GLEPP1	12	3.19	0.0870
948	208864_s_at	thioredoxin	Hs.76136.0	gb:AF313911.1		9	2.58	0.0873
949	200701_at	epididymal secretory protein (19.5kD)	Hs.119529.0	gb:NM_006432.1	HE1	14	2.84	0.0874
950	215157_x_at	poly(A)-binding protein, cytoplasmic 1	Hs.172182.2	gb:AI734929_RC		8	3.14	0.0874
951	213313_at	rab6 GTPase activating protein (GAP and centrosome-associated)	Hs.55099.1	gb:AI922519_RC		9	2.47	0.0874
952	209140_x_at	major histocompatibility complex, class I, B	Hs.77961.0	gb:L42024.1	HLA-B39	8	3.38	0.0875
953	212230_at	Homo sapiens phosphatidic acid phosphatase type 2B (PPAP2B), mRNA	Hs.173717.0	gb:AV725664		1	2.72	0.0877
954	202583_s_at	RAN binding protein 9	Hs.279886.0	gb:NM_005493.1	RANBP9	6	2.22	0.0878
955	201584_s_at	nuclear RNA helicase, DECD variant of DEAD box family	Hs.179606.0	gb:NM_005804.1	DDXL	19	2.28	0.0878
956	204020_at	purine-rich element binding protein A	Hs.29117.0	gb:BF739943_RC		5	2.51	0.0878
957	209150_s_at	transmembrane 9 superfamily member 1	Hs.91586.0	gb:U94831.1		14	2.11	0.0879
958	214721_x_at	Cdc42 effector protein 4; binder of Rho GTPases 4	Hs.3903.1	gb:AL162074.1	DKFZp76	17	2.22	0.0879
959	202527_s_at	MAD (mothers against decapentaplegic, Drosophila) homolog 4	Hs.75862.0	gb:NM_005359.1	MADH4	18	2.03	0.0883
960	210592_s_at	spermidinespermine N1-acetyltransferase	Hs.28491.1	gb:M55580.1	spermidin	10	3.03	0.0884
961	201865_x_at	nuclear receptor subfamily 3, group C, member 1	Hs.75772.0	gb:AI432196_RC		5	2.51	0.0884

Fig. 32

	A	B	C	D	E	F	G	H
962	214022_s_at	interferon induced transmembrane protein 1 (9-27)	Hs.146360.1	gb:AA749101_RC		12	2.90	0.0885
963	202854_at	hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome)	Hs.82314.0	gb:NM_000194.1	HPRT1	X	2.46	0.0886
964	201669_s_at	myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L)	Hs.75607.0	gb:NM_002356.4	MACS	6	2.63	0.0886
965	222243_s_at	transducer of ERBB2, 2	Hs.4994.1	gb:AB051450.1	KIAA166	22	2.04	0.0887
966	213151_s_at	CDC10 (cell division cycle 10, S. cerevisiae, homolog)	Hs.184326.1	gb:AU157515_RC		7	2.85	0.0887
967	200782_at	annexin A5	Hs.300711.0	gb:NM_001154.2	ANXA5	4	2.72	0.0889
968	201084_s_at	KIAA0164 gene product	Hs.80338.0	gb:NM_014739.1	KIAA016	6	2.60	0.0891
969	208636_at	actinin, alpha 1	Hs.119000.1	gb:AI082078_RC		14	2.69	0.0891
970	201177_s_at	SUMO-1 activating enzyme subunit 2	Hs.4311.0	gb:NM_005499.1	UBA2	19	2.25	0.0891
971	208091_s_at			gb:NM_030796.1	DKFZP56	7	2.47	0.0892
972	201426_s_at	vimentin	Hs.297753.0	gb:AI922599_RC		10	3.19	0.0893
973	201518_at	chromobox homolog 1 (Drosophila HP1 beta)	Hs.77254.0	gb:NM_006807.1	CBX1	17	2.29	0.0893
974	218603_at	hHDC for homolog of Drosophila headcase	Hs.6679.0	gb:NM_016217.1	LOC5169	6	2.32	0.0894
975	216308_x_at	glyoxylate reductasehydroxypyruvate reductase	Hs.155742.1	gb:AK026752.1		9	2.31	0.0894
976	200632_s_at	N-myc downstream regulated	Hs.75789.0	gb:NM_006096.1	NDRG1	8	2.76	0.0895
977	200595_s_at	eukaryotic translation initiation factor 3, subunit 10 (theta, 150170kD)	Hs.198899.0	gb:NM_003750.1	EIF3S10	10	2.45	0.0896
978	214039_s_at	Homo sapiens mRNA; cDNA DKFZp586E1124 (from clone DKFZp586E1	Hs.296398.1	gb:T15777_RC		3	2.83	0.0898
979	200641_s_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.75103.0	gb:U28964.1		8	2.63	0.0899
980	214512_s_at	activated RNA polymerase II transcription cofactor 4	Hs.74861.2	gb:NM_006713.1	PC4	8	2.31	0.0899
981	201938_at	deleted in oral cancer (mouse, homolog) 1	Hs.3436.0	gb:NM_004642.1	DOC1	12	2.76	0.0899
982	201296_s_at	DKFZP564A122 protein	Hs.187991.0	gb:NM_015626.1	DKFZP56	17	2.56	0.0902
983	200073_s_at	heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA-binding	Hs.303627.3	gb:M94630.1		4	2.32	0.0903
984	201285_at	makorin, ring finger protein, 1	Hs.7838.0	gb:NM_013446.1	MKRN1	7	2.51	0.0903
985	203832_at	small nuclear ribonucleoprotein polypeptide F	Hs.105465.0	gb:NM_003095.1	SNRPF	12	2.30	0.0904
986	201022_s_at	destrin (actin depolymerizing factor)	Hs.82306.0	gb:NM_006870.2	ADF	20	3.08	0.0905
987	200653_s_at	calmodulin 1 (phosphorylase kinase, delta)	Hs.177656.0	gb:M27319.1		14	2.86	0.0906
988	200961_at	selenophosphate synthetase 2	Hs.118725.0	gb:NM_012248.1	SPS2	5	2.28	0.0906
989	200593_s_at	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	Hs.103804.0	gb:BC003621.1		1	2.26	0.0908
990	215333_x_at	glutathione S-transferase M4	Hs.301961.2	gb:X08020.1		1	2.34	0.0908
991	201472_at	von Hippel-Lindau binding protein 1	Hs.198307.0	gb:NM_003372.2	VBP1	X	2.47	0.0909
992	217140_s_at	voltage-dependent anion channel 1 pseudogene	Hs.201553.0	gb:AJ002428			2.28	0.0910

Fig. 33

	A	B	C	D	E	F	G	H
993	201581_at	hypothetical protein	Hs.169358.0	gb:BF572868		20	2.33	0.0911
994	200708_at	glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransfe	Hs.170197.0	gb:NM_002080.1	GOT2	16	2.60	0.0911
995	203581_at	RAB4, member RAS oncogene family	Hs.119007.0	gb:BC002438.1		1	2.08	0.0911
996	200074_s_at	ribosomal protein L14	Hs.738.2	gb:U16738.1		12	2.89	0.0911
997	212612_at	KIAA0071 protein	Hs.78398.0	gb:D31888.1	KIAA007	14	1.96	0.0913
998	212470_at	sperm associated antigen 9	Hs.129872.1	gb:AB011088.1	KIAA051	17	2.24	0.0916
999	218478_s_at	hypothetical protein DKFZp434E2220	Hs.37706.0	gb:NM_017612.1	DKFZp43	12	2.04	0.0918
1000	208821_at	small nuclear ribonucleoprotein polypeptides B and B1	Hs.83753.0	gb:J04564.1	SNRPB	20	2.25	0.0918
1001	200902_at	15 kDa selenoprotein	Hs.90606.0	gb:NM_004261.1	15.Sep	1	2.45	0.0919
1002	212085_at	solute carrier family 25 (mitochondrial carrier; adenine nucleotide transloca	Hs.164280.1	gb:AA916851_RC		9	2.87	0.0919
1003	202169_s_at	aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferas	Hs.64595.0	gb:AF302110.1		11	2.41	0.0920
1004	208628_s_at	nuclease sensitive element binding protein 1	Hs.74497.1	gb:BC002411.1		1	3.02	0.0921
1005	205756_s_at	coagulation factor VIII, procoagulant component (hemophilia A)	Hs.79345.1	gb:NM_000132.2	F8	X	2.13	0.0922
1006	217717_s_at	GW128 protein	Hs.182238.0	gb:BF246499		20	2.48	0.0923
1007	201312_s_at	SH3 domain binding glutamic acid-rich protein like	Hs.14368.0	gb:NM_003022.1	SH3BGRI	X	2.62	0.0925
1008	205480_s_at	UDP-glucose pyrophosphorylase 2	Hs.77837.0	gb:NM_006759.2	UGP2		2.80	0.0926
1009	218450_at	heme-binding protein	Hs.108675.0	gb:NM_015987.1	HEBP	12	2.41	0.0928
1010	217356_s_at	Phosphoglycerate kinase {alternatively spliced} human, phosphoglycerate k	Hs.169313.0	gb:S81916.1		X	2.50	0.0930
1011	201652_at	COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5	Hs.198767.0	gb:NM_006837.1	COPS5	8	2.29	0.0930
1012	212341_at	Homo sapiens mRNA; cDNA DKFZp586F1822 (from clone DKFZp586F1	Hs.82719.0	gb:AA195936_RC			2.18	0.0933
1013	201360_at	cystatin C (amyloid angiopathy and cerebral hemorrhage)	Hs.135084.0	gb:NM_000099.1	CST3	20	3.21	0.0935
1014	200869_at	ribosomal protein L18a	Hs.163593.0	gb:NM_000980.1	RPL18A	19	3.27	0.0936
1015	205412_at	acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase)	Hs.37.0	gb:NM_000019.1	ACAT1	11	2.72	0.0937
1016	212010_s_at	hypothetical protein	Hs.283690.1	gb:AK025647.1		3	2.35	0.0938
1017	200931_s_at	vinculin	Hs.75350.0	gb:NM_014000.1	VCL	10	2.72	0.0939
1018	211962_s_at	butyrate response factor 1 (EGF-response factor 1)	Hs.85155.0	gb:BG250310		14	2.62	0.0939
1019	217725_x_at	PAI-1 mRNA-binding protein	Hs.165998.0	gb:NM_015640.1	PAI-RBP	1	2.48	0.0940
1020	201651_s_at	protein kinase C and casein kinase substrate in neurons 2	Hs.18842.0	gb:NM_007229.1	PACSIN2	22	2.63	0.0940
1021	202922_at	glutamate-cysteine ligase, catalytic subunit	Hs.151393.0	gb:BF676980		6	2.13	0.0941
1022	202076_at	baculoviral IAP repeat-containing 2	Hs.289107.0	gb:NM_001166.2	BIRC2	11	2.39	0.0942
1023	209409_at	growth factor receptor-bound protein 10	Hs.81875.3	gb:D86962.1	KIAA020	7	2.34	0.0942

Fig. 34

	A	B	C	D	E	F	G	H
1024	221829_s_at	karyopherin (importin) beta 2	Hs.168075.2	gb:AI307759_RC		5	2.59	0.0943
1025	212320_at	tubulin, beta polypeptide	Hs.179661.5	gb:BC001002.1		6	2.82	0.0943
1026	201041_s_at	dual specificity phosphatase 1	Hs.171695.0	gb:NM_004417.2	DUSP1	5	3.11	0.0945
1027	201064_s_at	poly(A)-binding protein, cytoplasmic 4 (inducible form)	Hs.169900.0	gb:NM_003819.2	PABPC4	1	2.56	0.0950
1028	212509_s_at	FK506 binding protein 12-rapamycin associated protein 1	Hs.250723.2	gb:BF968134		2	2.99	0.0953
1029	212412_at	Homo sapiens mRNA; cDNA DKFZp564A072 (from clone DKFZp564A0'	Hs.7780.0	gb:AV715767		4	2.57	0.0954
1030	214271_x_at	ribosomal protein L12	Hs.182979.2	gb:AA281332_RC		9	3.16	0.0956
1031	204034_at	protein expressed in thyroid	Hs.7486.0	gb:NM_014297.1	YF13H12	19	2.30	0.0957
1032	213574_s_at	karyopherin (importin) beta 1	Hs.180446.2	gb:AA861608_RC		X	2.40	0.0958
1033	218007_s_at	40S ribosomal protein S27 isoform	Hs.108957.0	gb:NM_015920.1	LOC5106	15	2.76	0.0958
1034	208857_s_at	protein-L-isoaspartate (D-aspartate) O-methyltransferase	Hs.79137.2	gb:M93008.1		6	2.41	0.0961
1035	212271_at	mitogen-activated protein kinase 1	Hs.324473.2	gb:AA195999_RC		22	2.31	0.0961
1036	214274_s_at	deleted in lung and esophageal cancer 1	Hs.200188.5	gb:AI860341_RC		3	2.08	0.0963
1037	201968_s_at	phosphoglucomutase 1	Hs.1869.0	gb:NM_002633.1	PGM1	1	2.70	0.0964
1038	200983_x_at	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5,	Hs.119663.0	gb:BF983379		11	2.88	0.0965
1039	201272_at	aldo-keto reductase family 1, member B1 (aldose reductase)	Hs.75313.0	gb:NM_001628.1	AKR1B1	7	2.69	0.0966
1040	201089_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypept	Hs.1697.0	gb:NM_001693.1	ATP6B2	8	2.57	0.0970
1041	213726_x_at	tubulin, beta, 2	Hs.251653.1	gb:AA515698_RC		X	2.90	0.0973
1042	200602_at	amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	Hs.177486.0	gb:NM_000484.1	APP	21	2.56	0.0976
1043	212276_at	lipin 1	Hs.81412.0	gb:D80010.1	KIAA018	2	2.31	0.0977
1044	208697_s_at	eukaryotic translation initiation factor 3, subunit 6 (48kD)	Hs.106673.0	gb:BC000734.1		8	2.84	0.0978
1045	214167_s_at	ribosomal protein, large, P0	Hs.73742.4	gb:AA555113_RC		12	3.04	0.0981
1046	200839_s_at	cathepsin B	Hs.297939.0	gb:NM_001908.1	CTSB	8	2.99	0.0985
1047	213274_s_at	cathepsin B	Hs.297939.5	gb:AA020826_RC		8	2.41	0.0987
1048	208953_at	KIAA0217 protein	Hs.78851.0	gb:AU154358_RC		10	1.92	0.0987
1049	214429_at	myotubularin related protein 6	Hs.79877.1	gb:U47635.1			2.01	0.0988
1050	209022_at	stromal antigen 2	Hs.8217.1	gb:AK026678.1		X	2.46	0.0989
1051	201627_s_at	insulin induced gene 1	Hs.56205.0	gb:NM_005542.1	INSIG1		2.27	0.0991
1052	200803_s_at	testis enhanced gene transcript (BAX inhibitor 1)	Hs.74637.0	gb:AF033095.1	TEGT	12	2.81	0.0994
1053	211404_s_at	amyloid beta (A4) precursor-like protein 2	Hs.279518.2	gb:BC004371.1		11	2.28	0.0994
1054	202365_at	acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain	Hs.127610.0	gb:BC004815.1		12	2.13	0.0996

Fig. 35

	A	B	C	D	E	F	G	H
1055	208977_x_at	tubulin, beta, 2	Hs.251653.0	gb:BC004188.1		6	2.89	0.0997
1056	201589_at	SMC1 (structural maintenance of chromosomes 1, yeast)-like 1	Hs.211602.0	gb:D80000.1	KIAA017	X	2.13	0.0999
1057	201737_s_at	similar to S. cerevisiae SSM4	Hs.20141.0	gb:NM_005885.1	TEB4	5	2.26	0.1003
1058	202606_s_at	tousled-like kinase 1	Hs.18895.0	gb:NM_012290.1	TLK1	2	2.07	0.1003
1059	201599_at	ornithine aminotransferase (gyrate atrophy)	Hs.75485.0	gb:NM_000274.1	OAT	10	2.62	0.1003
1060	200599_s_at	tumor rejection antigen (gp96) 1	Hs.82689.0	gb:NM_003299.1	TRA1	12	2.86	0.1004
1061	213168_at	Sp3 transcription factor	Hs.44450.1	gb:AU145005_RC		13	2.15	0.1004
1062	202727_s_at	interferon gamma receptor 1	Hs.180866.0	gb:NM_000416.1	IFNGR1	6	2.53	0.1006
1063	212638_s_at	Homo sapiens mRNA; cDNA DKFZp434D2111 (from clone DKFZp434D2	Hs.324275.0	gb:BF131791		3	2.28	0.1006
1064	200891_s_at	signal sequence receptor, alpha (translocon-associated protein alpha)	Hs.250773.0	gb:NM_003144.2	SSR1	6	2.54	0.1008
1065	212503_s_at	KIAA0934 protein	Hs.227716.0	gb:N22859_RC		10	2.29	0.1009
1066	210980_s_at	N-acylsphingosine amidohydrolase (acid ceramidase)	Hs.75811.1	gb:U47674.1		8	2.56	0.1010
1067	201004_at	signal sequence receptor, delta (translocon-associated protein delta)	Hs.102135.0	gb:NM_006280.1	SSR4	X	2.88	0.1010
1068	201503_at	Ras-GTPase-activating protein SH3-domain-binding protein	Hs.220689.0	gb:BG500067		5	2.11	0.1012
1069	201398_s_at	translocating chain-associating membrane protein	Hs.4147.0	gb:BC000687.1		8	2.63	0.1016
1070	202670_at	mitogen-activated protein kinase kinase 1	Hs.3446.0	gb:AI571419_RC		8	2.42	0.1017
1071	219126_at	hypothetical protein FLJ10975	Hs.321578.0	gb:NM_018288.1	FLJ10975	X	2.27	0.1018
1072	200994_at	RAN binding protein 7	Hs.5151.0	gb:BG291787		11	2.31	0.1022
1073	211733_x_at			gb:BC005911.1		1	2.37	0.1024
1074	210561_s_at	DKFZP564A122 protein	Hs.187991.2	gb:AL110243.1	DKFZp56	17	2.49	0.1026
1075	200823_x_at	ribosomal protein L29	Hs.183698.0	gb:NM_000992.1	RPL29	3	3.01	0.1027
1076	208991_at	signal transducer and activator of transcription 3 (acute-phase response fact	Hs.321677.0	gb:AA634272_RC		17	2.63	0.1028
1077	201522_x_at	small nuclear ribonucleoprotein polypeptide N	Hs.48375.4	gb:NM_003097.2	SNRPN	15	3.00	0.1033
1078	220948_s_at	ATPase, Na+K+ transporting, alpha 1 polypeptide	Hs.76549.0	gb:NM_000701.1	ATP1A1	1	2.79	0.1034
1079	203044_at	KIAA0990 protein	Hs.110488.0	gb:NM_014918.1	KIAA099	15	2.09	0.1040
1080	221676_s_at	coronin, actin-binding protein, 1C	Hs.17377.1	gb:BC002342.1		12	2.25	0.1041
1081	201318_s_at	myosin, light polypeptide, regulatory, non-sarcomeric (20kD)	Hs.233936.0	gb:NM_006471.1	MLCB	18	2.75	0.1044
1082	201991_s_at	kinesin family member 5B	Hs.149436.0	gb:BF223224_RC		10	2.54	0.1045
1083	203006_at	inositol polyphosphate-5-phosphatase, 40kD	Hs.124029.0	gb:NM_005539.1	INPP5A	10	2.46	0.1047
1084	214924_s_at	KIAA1042 protein	Hs.6705.1	gb:AK000754.1		3	2.16	0.1047
1085	216306_x_at	polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprot	Hs.172550.6	gb:X62006.1	PTB-1	NA	2.57	0.1049

Fig. 36

	A	B	C	D	E	F	G	H
1086	212825_at	PAX transcription activation domain interacting protein 1 like	Hs.173854.0	gb:AI357401_RC		7	1.88	0.1049
1087	200797_s_at	myeloid cell leukemia sequence 1 (BCL2-related)	Hs.86386.0	gb:AI275690_RC		1	2.81	0.1051
1088	200921_s_at	B-cell translocation gene 1, anti-proliferative	Hs.77054.0	gb:NM_001731.1	BTG1	12	2.69	0.1051
1089	211967_at	Homo sapiens mRNA; cDNA DKFZp586I2022 (from clone DKFZp586I20	Hs.172089.0	gb:BG538627		11	2.48	0.1057
1090	217853_at	hypothetical protein FLJ13732 similar to tensin	Hs.12210.0	gb:NM_022748.1	FLJ13732	7	2.46	0.1060
1091	208662_s_at	tetratricopeptide repeat domain 3	Hs.118174.2	gb:AI885338_RC		21	2.60	0.1061
1092	217745_s_at	hypothetical protein FLJ13194	Hs.288932.0	gb:NM_025146.1	FLJ13194	3	2.26	0.1061
1093	200698_at	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention recepto	Hs.118778.0	gb:AL542253_RC		7	2.50	0.1064
1094	200816_s_at	platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD)	Hs.77318.0	gb:NM_000430.2	PAFAH1I	17	2.50	0.1064
1095	214743_at	cut (Drosophila)-like 1 (CCAAT displacement protein)	Hs.147049.1	gb:BE046521_RC		7	2.18	0.1069
1096	202232_s_at	dendritic cell protein	Hs.69469.0	gb:NM_006360.1	GA17	11	2.23	0.1071
1097	203007_x_at	lysophospholipase I	Hs.12540.0	gb:AF077198.1		6	2.18	0.1073
1098	202749_at	tryptophan rich basic protein	Hs.198308.0	gb:NM_004627.1	WRB	21	2.37	0.1074
1099	202708_s_at	H2B histone family, member Q	Hs.2178.0	gb:NM_003528.1	H2BFQ	1	2.19	0.1074
1100	202459_s_at	lipin 2	Hs.166318.0	gb:U55968_RC		18	2.15	0.1074
1101	222035_s_at	poly(A) polymerase alpha	Hs.49007.1	gb:AI984479_RC		14	2.04	0.1079
1102	202166_s_at	protein phosphatase 1, regulatory (inhibitor) subunit 2	Hs.267819.0	gb:NM_006241.1	PPP1R2	3	2.11	0.1080
1103	208113_x_at			gb:NM_030979.1	LOC8187	13	2.81	0.1082
1104	243_g_at						2.32	0.1085
1105	205251_at	period (Drosophila) homolog 2	Hs.153405.0	gb:NM_022817.1	PER2	2	2.16	0.1085
1106	212762_s_at	transcription factor 7-like 2 (T-cell specific, HMG-box)	Hs.173638.0	gb:AI375916_RC		10	2.11	0.1085
1107	201432_at	catalase	Hs.76359.0	gb:NM_001752.1	CAT	11	2.82	0.1085
1108	217722_s_at	mesenchymal stem cell protein DSC92	Hs.323467.0	gb:NM_016645.1	LOC5133	15	2.79	0.1086
1109	203531_at	cullin 5	Hs.101299.0	gb:BF435809_RC		11	2.22	0.1090
1110	201010_s_at	upregulated by 1,25-dihydroxyvitamin D-3	Hs.179526.0	gb:NM_006472.1	VDUP1	1	3.12	0.1093
1111	208654_s_at	CD164 antigen, sialomucin	Hs.43910.2	gb:BF669455		6	2.52	0.1093
1112	205202_at	protein-L-isoaspartate (D-aspartate) O-methyltransferase	Hs.79137.0	gb:NM_005389.1	PCMT1	6	2.44	0.1097
1113	202754_at	KIAA0029 protein	Hs.268053.0	gb:NM_015361.1	KIAA002	2	2.25	0.1098
1114	213129_s_at	glycine cleavage system protein H (aminomethyl carrier)	Hs.77631.1	gb:AI970157_RC		16	2.47	0.1105
1115	201133_s_at	KIAA0438 gene product	Hs.279849.0	gb:AA142966_RC		5	2.23	0.1107
1116	212345_s_at	hypothetical protein DKFZp586F2423	Hs.13659.0	gb:BE675139_RC		7	2.40	0.1109

Fig. 37

	A	B	C	D	E	F	G	H
1117	208942_s_at	translocation protein 1	Hs.8146.0	gb:BE866511		3	2.44	0.1113
1118	39729_at						2.70	0.1115
1119	202804_at	ATP-binding cassette, sub-family C (CFTRMRP), member 1	Hs.89433.4	gb:AI539710_RC		16	2.12	0.1124
1120	203416_at	CD53 antigen	Hs.82212.0	gb:NM_000560.1	CD53	1	2.56	0.1126
1121	212181_s_at	nudix (nucleoside diphosphate linked moiety X)-type motif 4	Hs.92381.5	gb:AF191654.2		12	2.42	0.1127
1122	219549_s_at	reticulon 3	Hs.252831.0	gb:NM_006054.1	RTN3	4	2.65	0.1132
1123	209142_s_at	ubiquitin-conjugating enzyme E2G 1 (homologous to C. elegans UBC7)	Hs.78563.0	gb:BC002775.1		17	2.13	0.1136
1124	200700_s_at	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention recepto	Hs.118778.0	gb:NM_006854.2	KDELR2	7	2.63	0.1138
1125	200967_at	peptidylprolyl isomerase B (cyclophilin B)	Hs.699.0	gb:NM_000942.1	PPIB	15	2.55	0.1141
1126	200920_s_at	B-cell translocation gene 1, anti-proliferative	Hs.77054.0	gb:AL535380		12	2.81	0.1143
1127	210105_s_at	FYN oncogene related to SRC, FGR, YES	Hs.169370.0	gb:M14333.1		6	2.51	0.1145
1128	211945_s_at	Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antig	Hs.287797.0	gb:BG500301		10	2.88	0.1148
1129	220924_s_at	amino acid transporter 2	Hs.298275.0	gb:NM_018976.1	KIAA138	12	2.73	0.1148
1130	200634_at	profilin 1	Hs.75721.0	gb:NM_005022.1	PFN1	17	2.81	0.1151
1131	221731_x_at	chondroitin sulfate proteoglycan 2 (versican)	Hs.81800.4	gb:BF218922		5	2.40	0.1151
1132	202033_s_at	KIAA0203 gene product	Hs.50421.0	gb:BG402105		8	2.13	0.1152
1133	201931_at	electron-transfer-flavoprotein, alpha polypeptide (glutaric aciduria II)	Hs.169919.0	gb:NM_000126.1	ETFA	15	2.24	0.1154
1134	221253_s_at			gb:NM_030810.1	MGC317	6	2.63	0.1154
1135	210621_s_at	RAS p21 protein activator (GTPase activating protein) 1	Hs.758.1	gb:M23612.1	GAP	5	2.13	0.1161
1136	206042_x_at	SNRPN upstream reading frame	Hs.58606.0	gb:NM_022804.1	SNURF	15	2.45	0.1175
1137	201923_at	thioredoxin peroxidase (antioxidant enzyme)	Hs.83383.0	gb:NM_006406.1	AOE372	X	2.52	0.1179
1138	201020_at	tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein,	Hs.75544.0	gb:NM_003405.1	YWHAH	22	2.51	0.1179
1139	208937_s_at	inhibitor of DNA binding 1, dominant negative helix-loop-helix protein	Hs.75424.1	gb:D13889.1	Id-1H	20	2.62	0.1190
1140	200629_at	tryptophanyl-tRNA synthetase	Hs.82030.0	gb:NM_004184.2	WARS	14	2.36	0.1191
1141	201760_s_at	CS box-containing WD protein	Hs.136644.0	gb:NM_018639.1	LOC55884		2.50	0.1191
1142	218041_x_at	hypothetical protein PRO1068	Hs.321158.0	gb:NM_018573.1	PRO1068	12	2.74	0.1194
1143	217911_s_at	BCL2-associated athanogene 3	Hs.15259.0	gb:NM_004281.1	BAG3	10	2.44	0.1207
1144	202295_s_at	cathepsin H	Hs.288181.0	gb:NM_004390.1	CTSH	15	2.53	0.1208
1145	201590_x_at	annexin A2	Hs.217493.0	gb:NM_004039.1	ANXA2	15	3.05	0.1219
1146	200600_at	moesin	Hs.170328.0	gb:NM_002444.1	MSN	X	2.54	0.1225
1147	208637_x_at	actinin, alpha 1	Hs.119000.1	gb:BC003576.1		14	2.42	0.1228

Fig. 38

	A	B	C	D	E	F	G	H
1148	201339_s_at	sterol carrier protein 2	Hs.75760.0	gb:NM_002979.1	SCP2	1	2.47	0.1236
1149	201540_at	four and a half LIM domains 1	Hs.239069.0	gb:NM_001449.1	FHL1	X	3.02	0.1238
1150	219017_at	ethanolamine kinase	Hs.120439.0	gb:NM_018638.2	EKI1	12	1.85	0.1258
1151	201088_at	karyopherin alpha 2 (RAG cohort 1, importin alpha 1)	Hs.159557.0	gb:NM_002266.1	KPNA2	17	2.28	0.1259
1152	218205_s_at	G protein-coupled receptor kinase 7	Hs.261828.0	gb:NM_017572.1	GPRK7	NA	2.62	0.1265
1153	211458_s_at	Homo sapiens GABA-A receptor-associated protein mRNA, complete cds	Hs.326776.0	gb:AF180519.1		15	2.59	0.1266
1154	201963_at	fatty-acid-Coenzyme A ligase, long-chain 2	Hs.154890.0	gb:NM_021122.2	FACL2	4	2.48	0.1282
1155	200733_s_at	protein tyrosine phosphatase type IVA, member 1	Hs.227777.0	gb:U48296.1	hPTPCAP	6	1.97	0.1287
1156	219032_x_at	opsin 3 (encephalopsin)	Hs.279926.0	gb:NM_014322.1	OPN3	1	2.09	0.1290
1157	217908_s_at	PC326 protein	Hs.279882.0	gb:NM_018442.1	PC326	1	2.05	0.1295
1158	217979_at	tetraspan NET-6 protein	Hs.284243.0	gb:NM_014399.1	NET-6	7	2.55	0.1310
1159	202581_at	heat shock 70kD protein 1B	Hs.274402.0	gb:NM_005346.2	HSPA1B	6	2.39	0.1323
1160	201486_at	reticulocalbin 2, EF-hand calcium binding domain	Hs.79088.0	gb:NM_002902.1	RCN2	15	2.12	0.1326
1161	205353_s_at	prostatic binding protein	Hs.80423.0	gb:NM_002567.1	PBP	2	2.75	0.1327
1162	208623_s_at	villin 2 (ezrin)	Hs.155191.0	gb:J05021.1	VIL2	6	2.34	0.1333
1163	212501_at	CCAATenhancer binding protein (CEBP), beta	Hs.99029.0	gb:AL564683_RC		20	2.87	0.1341
1164	221589_s_at	methylmalonate-semialdehyde dehydrogenase	Hs.293970.1	gb:AW612403_RC		14	2.22	0.1348
1165	201864_at	GDP dissociation inhibitor 1	Hs.74576.0	gb:NM_001493.1	GDI1	X	2.67	0.1350
1166	200788_s_at	phosphoprotein enriched in astrocytes 15	Hs.194673.0	gb:NM_003768.1	PEA15	1	2.79	0.1371
1167	212928_at	KIAA0721 protein	Hs.284141.0	gb:AL050331		6	2.44	0.1380
1168	221011_s_at			gb:NM_030915.1	DKFZP56	2	2.25	0.1386
1169	201348_at	glutathione peroxidase 3 (plasma)	Hs.172153.0	gb:NM_002084.2	GPX3	5	3.01	0.1394
1170	209340_at	UDP-N-acteylglucosamine pyrophosphorylase 1	Hs.21293.0	gb:S73498.1		1	2.16	0.1396
1171	212086_x_at	lamin AC	Hs.77886.2	gb:AK026584.1		1	2.27	0.1400
1172	200644_at	macrophage myristoylated alanine-rich C kinase substrate	Hs.75061.0	gb:NM_023009.1	MACMAI	1	2.75	0.1406
1173	203335_at	phytanoyl-CoA hydroxylase (Refsum disease)	Hs.172887.0	gb:NM_006214.1	PHYH	10	2.24	0.1412
1174	208691_at	transferrin receptor (p90, CD71)	Hs.77356.1	gb:BC001188.1		3	2.57	0.1442
1175	208944_at	transforming growth factor, beta receptor II (70-80kD)	Hs.82028.1	gb:D50683.1		3	2.43	0.1449
1176	201242_s_at	ATPase, Na+K+ transporting, beta 1 polypeptide	Hs.78629.0	gb:BC000006.1		1	2.94	0.1470
1177	215772_x_at	succinate-CoA ligase, GDP-forming, beta subunit	Hs.247309.2	gb:AL050226.1	DKFZp58	12	2.23	0.1481
1178	212459_x_at	succinate-CoA ligase, GDP-forming, beta subunit	Hs.247309.0	gb:BF593940_RC		3	2.24	0.1503

Fig. 39

	A	B	C	D	E	F	G	H
1179	208892_s_at	dual specificity phosphatase 6	Hs.180383.1	gb:BC003143.1		12	2.11	0.1506
1180	212508_at	modulator of apoptosis 1	Hs.24719.0	gb:AK024029.1		14	2.42	0.1532
1181	214091_s_at	glutathione peroxidase 3 (plasma)	Hs.172153.1	gb:AW149846_RC		5	2.81	0.1583
1182	212774_at	zinc finger protein 238	Hs.69997.1	gb:AJ223321		1	2.41	0.1610
1183	209118_s_at	Tubulin, alpha, brain-specific	Hs.272897.0	gb:AF141347.1		12	2.92	0.1620
1184	211985_s_at	matrix Gla protein	Hs.279009.6	gb:AI653730_RC		14	2.66	0.1621
1185	210986_s_at	tropomyosin 1 (alpha)	Hs.77899.1	gb:Z24727.1		15	2.66	0.1651
1186	200670_at	X-box binding protein 1	Hs.149923.0	gb:NM_005080.1	XBP1	22	2.58	0.1657
1187	201160_s_at	cold shock domain protein A	Hs.1139.0	gb:AL556190		12	2.84	0.1665
1188	211896_s_at	decorin	Hs.76152.3	gb:AF138302.1		12	2.55	0.2165
1189	200606_at	desmoplakin (DPI, DPII)	Hs.74316.0	gb:NM_004415.1	DSP	6	2.45	0.2169

Fig. 40

No.	Tissue	Source		Lot No.		Sex	Age (years)	Race
1	Heart	Ambion7966	Total RNA	110P43B	Single	Male	56	Caucasian
2	Thymus	Ambion 7964	Total RNA	101P0101A	Single	Male	14	Caucasian
3	Spleen	Ambion 7970	Total RNA	061P18A	Single	Male	25	Caucasian
4	Ovary	Ambion 7974	Total RNA	051P42A	Single	Female	47	Caucasian
5	Kidney	Ambion 7976	Total RNA	071P04B	Single	Male	66	Asian
6	Skeltal Muscle	Ambion 7982	Total RNA	091P0101C	Single	Female	44	Caucasian
7	Pancreas	Ambion 7954	Total RNA	091P0104A	Single	Female	68	Caucasian
8	Prostate	Ambion 7988	Total RNA	081P0103A	Single	Male	26	Caucasian
9	Small Instesine	Ambion 7984	Total RNA	091P0201A	Single	Male	85	Caucasian
10	Colon	Ambion 7986	Total RNA	071P10B	Single	Female	34	Caucasian
11	Placenta	Ambion 7950	Total RNA	061P33B	Single	Female	30	Black
12	Bladder	Ambion 7990	Total RNA	81P0101A	Single	Male	28	Caucasian
13	Breast	Strategene 735044	Total RNA	0610327	Single	Female	44	
14	Uterus	Strategene 735042	Total RNA	1100640	Pooled	Female	54, 68, 76	
15	Thyroid	Strategene 735040	Total RNA	0510225	Single	Female	42	
16	Skin	Strategene 735031	Total RNA	0120484	Pool of 2	Male	18 and 19	
17	Salivary gland	Clontech 64026-1	Total RNA	1011322	Pooled	male/female	15-60	Caucasians
18	Trachea	Clontech 64091-1	Total RNA	1010201	Pool of 84	male/female	17-70	Caucasians
19	Brain, Cerebellum	Clontech 64035-1	Total RNA	1010033	Single	Male	25	Asian
20	Brain, Whole	Clontech 64020-1	Total RNA	0101041	Single	Male	28	Asian
21	Fetal brain	Clontech64094-1	Total RNA	2020902	Pool of 59	male/female	20-33weeks	Caucasians
22	Adrenal gland	Clontech64096-1	Total RNA	2020671	Pool of 61	male/female	15-61	Caucasians
23	Bone Marrow	Clontech64106-1	Total RNA	1110932	Pool of 8	male/female	22-65	Caucasians
24	amygdala	Clontech 6574-1	PolyA RNA	1030830	Pool of 70	male/female	17-76	Caucasians
25	caudate	Clontech 6575-1	PolyA RNA	0120289	Pool of 57	male/female	18-78	Caucasians
26	corpus callosum	Clontech 6577-1	PolyA RNA	1010486	Pool of 47	male/female	19-76	Caucasians
27	hippocampus	Clontech 6578-1	PolyA RNA	1050638	Pool of 19	male/female	16-70	Caucasians
28	thalamus	Clontech 6582-1	PolyA RNA	1070147	Pool of 10	male/female	32-75	Caucasians
29	pituitary gland	Clontech 6584-1	PolyA RNA	2010981	Pool of 12,	male/female	18-52	Caucasians
30	spinal cord	Clontech 6593-1	PolyA RNA	0111062	Pool of 49	male/female	15-66	Caucasians
31	Testis	Clontech64027-1	Total RNA	6120257	Pool of 19	Male	17 - 61	Caucasians
32	Liver	Internal	Surgery	N4	Single	Male	66	Japanese
33	Stomach	Internal	Surgery	MN15	Single	Male	77	Japanese
34	Lung	Internal	Surgery	14887	Single	Male	70	Japanese
35	Fetal liver	CHEMICON356	Total RNA	21060678	Pool of at least 3			

Fig. 41

	A	B	C	D	E	F	G	H	I	J	K	L
1	Whole Brain	205358_at	glutamate receptor, ionotropic, AMPA 2	Hs.89582.0	gb:NM_000826.1	GRIA2	4	14	549	2.74	0.31	0.17
2	Whole Brain	203999_at	synaptotagmin I	Hs.154679.0	gb:AV731490		12	13.7	2861	3.46	1.93	0.11
3	Whole Brain	216617_s_at	myelin associated glycoprotein	Hs.1780.1	gb:X98405.1	S-MAG	19	13.4	129	2.11	0.27	0.14
4	Whole Brain	214762_at	Homo sapiens DNA, cosmid clones TN62 and TN82	Hs.249227.0	gb:BF340635		6	12.2	1110	3.05	1.68	0.11
5	Whole Brain	221236_s_at			gb:NM_030795.1	RB3	8	11.9	1015	3.01	1.65	0.11
6	Whole Brain	206384_at	calcium channel, voltage-dependent, gamma subur	Hs.7235.0	gb:NM_006539.1	CACNG3	16	11.2	206	2.31	0.66	0.15
7	Whole Brain	213040_s_at	neuronal pentraxin receptor	Hs.91622.0	gb:AL008583		22	11	284	2.45	0.56	0.17
8	Whole Brain	206013_s_at	actin-like 6	Hs.259831.0	gb:NM_016188.2	ACTL6	7	10.7	153	2.18	0.22	0.18
9	Whole Brain	205031_at	ephrin-B3	Hs.26988.0	gb:NM_001406.1	EFNB3	17	10.5	455	2.66	1.25	0.13
10	Whole Brain	205143_at	chondroitin sulfate proteoglycan 3 (neurocan)	Hs.169047.0	gb:NM_004386.1	CSPG3	19	10.3	586	2.77	1.55	0.12
11	Whole Brain	211679_x_at			gb:AF095784.1	GABBR2	9	10.2	708	2.85	0.97	0.18
12	Whole Brain	207323_s_at	myelin basic protein	Hs.69547.0	gb:NM_002385.1	MBP	18	9.9	2351	3.37	1.48	0.19
13	Whole Brain	208321_s_at	calcium binding protein 1 (calbrain)	Hs.301423.0	gb:NM_004276.1	CABP1		9.8	624	2.80	0.38	0.25
14	Whole Brain	213676_at	Homo sapiens germline mRNA sequence	Hs.12840.0	gb:AL038824_RC		6	9.8	635	2.80	1.67	0.12
15	Whole Brain	213841_at	Homo sapiens mRNA, chromosome 1 specific transc	Hs.301281.0	gb:BE223030_RC		1	9.7	531	2.73	0.93	0.19
16	Whole Brain	206678_at	gamma-aminobutyric acid (GABA) A receptor, alpl	Hs.45740.0	gb:NM_000806.2	GABRA1	5	9.5	284	2.45	0.87	0.17
17	Whole Brain	209871_s_at	amyloid beta (A4) precursor protein-binding, fa	Hs.26468.0	gb:AB014719.1	hX11L	15	9.2	527	2.72	0.89	0.20
18	Whole Brain	210016_at	KIAA1106 protein	Hs.172619.0	gb:BF223003_RC		2	9.1	549	2.74	1.35	0.15
19	Whole Brain	209556_at	neurochondrin	Hs.94653.1	gb:AB011179.1	KIAA0607	1	9	692	2.84	1.72	0.12
20	Whole Brain	204338_s_at	regulator of G-protein signalling 4	Hs.227571.0	gb:NM_005613.2	RGS4	1	8.9	182	2.26	0.30	0.22
21	Whole Brain	209839_at	KIAA0820 protein	Hs.33578.0	gb:AL136712.1	DKFZp566K0	1	8.8	592	2.77	1.58	0.14
22	Whole Brain	209197_at	KIAA0080 protein	Hs.74554.0	gb:AA626780_RC		1	8.7	1243	3.09	1.74	0.16
23	Whole Brain	220131_at	FXYD domain-containing ion transport regulator	Hs.134729.0	gb:NM_022006.1	FXYD7	19	8.6	464	2.67	1.57	0.13
24	Whole Brain	205454_at	hippocalcin	Hs.288654.0	gb:BC001777.1		1	8.3	563	2.75	0.51	0.27
25	Whole Brain	206196_s_at	RaP2 interacting protein 8	Hs.6755.0	gb:NM_006695.1	RPIP8	17	8.3	1271	3.10	1.73	0.17
26	Whole Brain	215161_at	similar to calciumcalmodulin dependent protein	Hs.199068.0	gb:AW016039_RC		1	8.3	137	2.14	0.93	0.15
27	Whole Brain	221916_at	hypothetical protein	Hs.302689.1	gb:BF055311_RC		8	8.3	1473	3.17	1.60	0.19
28	Whole Brain	39966_at						8	1076	3.03	1.35	0.21
29	Whole Brain	204229_at	brain-specific Na-dependent inorganic phosphate	Hs.6535.0	gb:H40895_RC		19	7.8	1374	3.14	1.72	0.18
30	Whole Brain	219425_at	sulfortranferase family 4A, member 1	Hs.189810.0	gb:NM_014351.1	SULTX3	22	7.5	843	2.93	1.50	0.19
31	Whole Brain	209990_s_at	G protein-coupled receptor 51	Hs.198612.0	gb:AF056085.1		9	7.4	1346	3.13	1.23	0.26
32	Whole Brain	213601_at	slit (Drosophila) homolog 1	Hs.133466.1	gb:AB011537.2	MEGF4	10	7.4	725	2.86	1.89	0.13
33	Whole Brain	204586_at	bassoon (presynaptic cytomatrix protein)	Hs.194684.0	gb:NM_003458.1	BSN	NA	7.2	319	2.50	0.95	0.22
34	Whole Brain	204762_s_at	guanine nucleotide binding protein (G protein),	Hs.296184.0	gb:BE670563_RC		16	7.2	253	2.40	1.69	0.10
35	Whole Brain	208320_at	calcium binding protein 1 (calbrain)	Hs.301423.0	gb:NM_004276.1	CABP1		7.2	375	2.57	0.27	0.32
36	Whole Brain	204763_s_at	guanine nucleotide binding protein (G protein),	Hs.296184.0	gb:NM_020988.1	GNAO1	16	7.1	620	2.79	1.28	0.21
37	Whole Brain	209462_at	amyloid beta (A4) precursor-like protein 1	Hs.74565.0	gb:U48437.1		19	7.1	964	2.98	0.70	0.32
38	Whole Brain	205737_at	potassium voltage-gated channel, KQT-like subfa	Hs.4975.0	gb:NM_004518.1	KCNQ2	20	6.9	477	2.68	1.40	0.19

Fig. 42

	A	B	C	D	E	F	G	H	I	J	K	L
39	Whole Brain	206089_at	nel (chicken)-like 1	Hs. 21602. 0	gb:NM_006157. 1	NELL1	11	6. 7	151	2. 18	0. 94	0. 19
40	Whole Brain	206349_at	leucine-rich, glioma inactivated 1	Hs. 194704. 0	gb:NM_005097. 1	LGI1	10	6. 7	89	1. 95	0. 51	0. 22
41	Whole Brain	206773_at	lymphocyte antigen 6 complex, locus H	Hs. 159590. 0	gb:NM_002347. 1	LY6H		6. 7	297	2. 47	1. 34	0. 17
42	Whole Brain	219671_at	hippocalcin-like protein 4	Hs. 272891. 0	gb:AL136591. 1	DKFZp761G1	1	6. 6	519	2. 72	1. 69	0. 16
43	Whole Brain	203540_at	glial fibrillary acidic protein	Hs. 1447. 0	gb:NM_002055. 1	GFAP	17	6. 5	1987	3. 30	1. 37	0. 30
44	Whole Brain	213411_at	Homo sapiens mRNA; cDNA DKFZp434E0528 (from cl	Hs. 256398. 0	gb:AW242701_RC		7	6. 5	360	2. 56	1. 47	0. 17
45	Whole Brain	215005_at	hypothetical protein	Hs. 140950. 0	gb:AV723666		16	6. 4	175	2. 24	0. 97	0. 20
46	Whole Brain	216963_s_at	growth associated protein 43	Hs. 79000. 1	gb:AF279774. 1		3	6. 4	609	2. 78	1. 82	0. 15
47	Whole Brain	219236_at	hypothetical protein FLJ22672	Hs. 235873. 0	gb:NM_024897. 1	FLJ22672	1	6. 4	1396	3. 14	1. 65	0. 23
48	Whole Brain	203618_at	lifeguard	Hs. 182859. 0	gb:AB023167. 1	KIAA0950	12	6. 3	419	2. 62	1. 83	0. 13
49	Whole Brain	204730_at	KIAA0237 gene product	Hs. 78748. 0	gb:NM_014747. 1	KIAA0237	1	6. 3	1044	3. 02	1. 58	0. 23
50	Whole Brain	207010_at	gamma-aminobutyric acid (GABA) A receptor, beta	Hs. 89768. 0	gb:NM_000812. 2	GABRB1	4	6. 3	251	2. 40	1. 38	0. 16
51	Whole Brain	213482_at	dedicator of cyto-kinesis 3	Hs. 7022. 0	gb:BF593175_RC		3	6. 3	214	2. 33	0. 63	0. 27
52	Whole Brain	219415_at	tweety (Drosophila) homolog 1	Hs. 12828. 0	gb:NM_020659. 1	TTYH1	19	6. 3	311	2. 49	0. 58	0. 31
53	Whole Brain	220889_s_at	Carbonic anhydrase-related protein 10	Hs. 283110. 0	gb:NM_020178. 1	LOC56934	17	6. 3	114	2. 06	0. 42	0. 26
54	Whole Brain	206448_at	KIAA0844 protein	Hs. 22653. 0	gb:NM_014951. 1	KIAA0844	10	6. 2	709	2. 85	1. 46	0. 22
55	Whole Brain	213276_at	Homo sapiens clone 25048 mRNA sequence	Hs. 321572. 0	gb:T15766_RC		7	6. 2	337	2. 53	0. 74	0. 29
56	Whole Brain	214230_at	cell division cycle 42 (GTP-binding protein, 25	Hs. 146409. 3	gb:R37664_RC		1	6. 2	175	2. 24	1. 35	0. 15
57	Whole Brain	217077_s_at	G protein-coupled receptor 51	Hs. 198612. 1	gb:AF095723. 1		9	6. 2	475	2. 68	0. 89	0. 29
58	Whole Brain	218623_at	HMP19 protein	Hs. 70669. 0	gb:NM_015980. 1	LOC51617	5	6. 2	822	2. 91	0. 98	0. 31
59	Whole Brain	205818_at	deleted in bladder cancer chromosome region can	Hs. 6090. 0	gb:NM_014618. 1	DBCCR1	9	6. 1	331	2. 52	1. 73	0. 13
60	Whole Brain	214650_x_at	myelin oligodendrocyte glycoprotein	Hs. 141308. 2	gb:AL050328		6	6. 1	428	2. 63	1. 52	0. 18
61	Whole Brain	220359_s_at	cyclic AMP-regulated phosphoprotein, 21 kD	Hs. 7459. 0	gb:NM_016300. 1	ARPP-21	3	6. 1	490	2. 69	0. 82	0. 31
62	Whole Brain	203488_at	lectomedin-2	Hs. 107054. 0	gb:NM_014921. 1	KIAA0821	19	6	224	2. 35	1. 72	0. 10
63	Whole Brain	210040_at	solute carrier family 12, (potassium-chloride t	Hs. 21413. 0	gb:AF208159. 1	KCC2	20	6	994	3. 00	1. 50	0. 25
64	Whole Brain	214376_at	Homo sapiens clone 24626 mRNA sequence	Hs. 13438. 0	gb:AI263044_RC		4	6	219	2. 34	1. 49	0. 14
65	Whole Brain	215116_s_at	dynamin 1	Hs. 166161. 2	gb:AF035321. 1		9	5. 9	1537	3. 19	1. 66	0. 26
66	Whole Brain	221214_s_at	DKFZP586J1624 protein	Hs. 4964. 0	gb:NM_015537. 1	DKFZP586J1624		5. 9	557	2. 75	1. 89	0. 15
67	Whole Brain	206888_s_at	Rho GDP dissociation inhibitor (GDI) gamma	Hs. 121516. 0	gb:U82532. 1		16	5. 8	445	2. 65	1. 60	0. 18
68	Whole Brain	214933_at	calcium channel, voltage-dependent, PQ type, a1	Hs. 96253. 2	gb:AA769818_RC		19	5. 8	267	2. 43	1. 78	0. 11
69	Whole Brain	222020_s_at	neurotrimin	Hs. 288433. 2	gb:AW117456_RC		11	5. 8	126	2. 10	0. 63	0. 25
70	Whole Brain	206731_at	KIAA0902 protein	Hs. 100527. 0	gb:NM_014927. 1	KIAA0902	X	5. 7	206	2. 31	1. 29	0. 18
71	Whole Brain	214822_at	hypothetical protein from clone 24828	Hs. 21708. 0	gb:AF131833. 1		1	5. 7	199	2. 30	1. 02	0. 22
72	Whole Brain	219896_at	calcyon; D1 dopamine receptor-interacting prote	Hs. 148680. 0	gb:NM_015722. 2	CALCYON	10	5. 7	402	2. 60	1. 31	0. 23
73	Whole Brain	220331_at	cytochrome P450, subfamily 46 (cholesterol 24-h	Hs. 25121. 0	gb:NM_006668. 1	CYP46	14	5. 7	969	2. 99	1. 43	0. 27
74	Whole Brain	202517_at	collapsin response mediator protein 1	Hs. 155392. 0	gb:NM_001313. 1	CRMP1	4	5. 6	627	2. 80	1. 80	0. 18
75	Whole Brain	203389_at	kinesin family member 3C	Hs. 21611. 0	gb:AF035621. 1	KIF3C	2	5. 6	592	2. 77	1. 70	0. 19
76	Whole Brain	205635_at	huntingtin-associated protein interacting prote	Hs. 8004. 0	gb:NM_003947. 1	HAPIP	3	5. 6	440	2. 64	1. 02	0. 29

Fig. 43

	A	B	C	D	E	F	G	H	I	J	K	L
77	Whole Brain	205970_at	metallothionein 3 (growth inhibitory factor (ne	Hs. 73133. 0	gb:NM_005954. 1	MT3	16	5. 6	1939	3. 29	1. 64	0. 30
78	Whole Brain	207853_s_at	synuclein, beta	Hs. 90297. 0	gb:NM_003085. 2	SNCB	5	5. 6	483	2. 68	1. 02	0. 30
79	Whole Brain	209000_s_at	KIAA0202 protein	Hs. 80712. 0	gb:BC001329. 1		5	5. 6	474	2. 68	0. 71	0. 35
80	Whole Brain	213496_at	KIAA0455 gene product	Hs. 13245. 0	gb:AW592563_RC		1	5. 6	339	2. 53	1. 52	0. 18
81	Whole Brain	214111_at	Homo sapiens clone 24461 mRNA sequence	Hs. 4817. 0	gb:AF070577. 1		11	5. 6	490	2. 69	0. 53	0. 39
82	Whole Brain	221801_x_at	neurofilament, light polypeptide (68kD)	Hs. 211584. 0	gb:AL566528_RC		8	5. 6	177	2. 25	0. 44	0. 32
83	Whole Brain	222005_s_at	guanine nucleotide binding protein (G protein),	Hs. 179915. 1	gb:AL538966		11	5. 6	1011	3. 00	1. 87	0. 20
84	Whole Brain	202508_s_at	synaptosomal-associated protein, 25kD	Hs. 84389. 0	gb:NM_003081. 1	SNAP25	20	5. 5	2065	3. 31	1. 34	0. 36
85	Whole Brain	209470_s_at	glycoprotein M6A	Hs. 75819. 0	gb:D49958. 1		4	5. 5	1444	3. 16	1. 49	0. 30
86	Whole Brain	212880_at	WD repeat domain 7	Hs. 10881. 0	gb:AB011113. 1	KIAA0541	18	5. 5	339	2. 53	1. 97	0. 10
87	Whole Brain	220115_s_at	cadherin 10, type 2 (T2-cadherin)	Hs. 92489. 0	gb:NM_006727. 1	CDH10	5	5. 5	209	2. 32	1. 65	0. 12
88	Whole Brain	205230_at	KIAA0985 protein	Hs. 21239. 0	gb:NM_014954. 1	KIAA0985	12	5. 4	804	2. 91	1. 47	0. 27
89	Whole Brain	205279_s_at	glycine receptor, beta	Hs. 32973. 0	gb:AF094754. 1	GLRB	4	5. 4	249	2. 40	1. 68	0. 13
90	Whole Brain	206330_s_at	neuronal Shc	Hs. 151123. 0	gb:NM_016848. 1	SHC3	9	5. 4	265	2. 42	0. 77	0. 31
91	Whole Brain	206397_x_at	growth differentiation factor 1	Hs. 92614. 0	gb:NM_001492. 3	GDF1	19	5. 4	370	2. 57	0. 94	0. 30
92	Whole Brain	213200_at	synaptophysin	Hs. 75667. 0	gb:U93305		X	5. 4	299	2. 48	1. 45	0. 19
93	Whole Brain	64488_at						5. 3	168	2. 22	1. 68	0. 10
94	Whole Brain	205551_at	KIAA0735 gene product; synaptic vesicle protein	Hs. 8071. 0	gb:NM_014848. 1	KIAA0735	15	5. 3	1133	3. 05	1. 20	0. 35
95	Whole Brain	207093_s_at	oligodendrocyte myelin glycoprotein	Hs. 194772. 0	gb:NM_002544. 1	OMG	17	5. 3	826	2. 92	1. 82	0. 21
96	Whole Brain	214293_at	hypothetical protein FLJ10849	Hs. 8768. 1	gb:AI539361_RC		4	5. 3	241	2. 38	1. 62	0. 14
97	Whole Brain	219170_at	hypothetical protein MGC3213	Hs. 28144. 0	gb:NM_024333. 1	MGC3213	19	5. 3	295	2. 47	0. 97	0. 28
98	Whole Brain	39650_s_at						5. 2	124	2. 09	1. 58	0. 10
99	Whole Brain	209794_at	KIAA0411 gene product	Hs. 7977. 0	gb:AB007871. 1	KIAA0411	3	5. 2	459	2. 66	1. 76	0. 17
100	Whole Brain	222091_at	hippocalcin-like protein 4	Hs. 272891. 1	gb:BE550384_RC		1	5. 2	96	1. 98	0. 76	0. 23
101	Whole Brain	205278_at	glutamate decarboxylase 1 (brain, 67kD)	Hs. 324784. 1	gb:NM_000817. 1	GAD1	2	5. 1	651	2. 81	0. 84	0. 39
102	Whole Brain	205368_at	KIAA0773 gene product	Hs. 135343. 0	gb:NM_014690. 1	KIAA0773		5. 1	239	2. 38	1. 11	0. 25
103	Whole Brain	213484_at	Homo sapiens clone 23700 mRNA sequence	Hs. 66187. 0	gb:AI097640_RC		2	5. 1	488	2. 69	1. 14	0. 31
104	Whole Brain	215267_s_at	KIAA1087 protein	Hs. 143026. 0	gb:AI127885_RC		19	5. 1	208	2. 32	1. 00	0. 26
105	Whole Brain	204720_s_at	DnaJ (Hsp40) homolog, subfamily B, member 6	Hs. 44896. 0	gb:AV729634		1	5	949	2. 98	1. 37	0. 32
106	Whole Brain	204993_at	guanine nucleotide binding protein (G protein),	Hs. 92002. 0	gb:NM_002073. 1	GNAZ	22	5	343	2. 54	1. 88	0. 13
107	Whole Brain	205814_at	glutamate receptor, metabotropic 3	Hs. 3786. 0	gb:NM_000840. 1	GRM3	7	5	289	2. 46	1. 25	0. 24
108	Whole Brain	206179_s_at	brain-specific protein p25 alpha	Hs. 29353. 0	gb:NM_007030. 1	p25	5	5	104	2. 02	0. 79	0. 25
109	Whole Brain	209198_s_at	KIAA0080 protein	Hs. 74554. 0	gb:BC004291. 1		1	5	123	2. 09	1. 41	0. 14
110	Whole Brain	210181_s_at	calcium binding protein 1 (calbrain)	Hs. 301423. 1	gb:AF169148. 1	CABP1	12	5	369	2. 57	0. 65	0. 39
111	Whole Brain	210185_at	calcium channel, voltage-dependent, beta 1 subu	Hs. 635. 2	gb:AB054985. 1		17	5	187	2. 27	1. 79	0. 09
112	Whole Brain	211483_x_at	calciumcalmodulin-dependent protein kinase (CaM	Hs. 4884. 5	gb:AF081924. 1	CAMKB		5	797	2. 90	1. 62	0. 26
113	Whole Brain	219737_s_at	protocadherin 9	Hs. 12450. 0	gb:AI524125_RC		13	5	586	2. 77	1. 70	0. 21
114	Whole Brain	44040_at						4. 9	468	2. 67	1. 83	0. 17

Fig. 44

	A	B	C	D	E	F	G	H	I	J	K	L
115	Whole Brain	203069_at	KIAA0736 gene product	Hs. 7979. 0	gb:NM_014849. 1	KIAA0736	1	4. 9	747	2. 87	0. 94	0. 39
116	Whole Brain	205257_s_at	amphiphysin (Stiff-Mann syndrome with breast ca	Hs. 173034. 0	gb:NM_001635. 1	AMPH	7	4. 9	423	2. 63	1. 60	0. 21
117	Whole Brain	206135_at	KIAA0535 gene product	Hs. 151449. 0	gb:NM_014682. 1	KIAA0535	8	4. 9	50	1. 70	0. 26	0. 29
118	Whole Brain	206935_at	protocadherin 8	Hs. 19492. 0	gb:NM_002590. 2	PCDH8	13	4. 9	241	2. 38	0. 82	0. 32
119	Whole Brain	213558_at	piccolo (presynaptic cytomatrix protein)	Hs. 12376. 1	gb:AB011131. 1	KIAA0559	7	4. 9	232	2. 37	1. 65	0. 15
120	Whole Brain	219619_at	hypothetical protein DKFZp761C07121	Hs. 165636. 0	gb:NM_017594. 1	DKFZp761C0	9	4. 9	682	2. 83	1. 15	0. 35
121	Whole Brain	203619_s_at	lifeguard	Hs. 182859. 0	gb:NM_012306. 1	KIAA0950	12	4. 8	849	2. 93	1. 07	0. 39
122	Whole Brain	208457_at	gamma-aminobutyric acid (GABA) A receptor, del	Hs. 113882. 0	gb:NM_000815. 1	GABRD	1	4. 8	124	2. 09	0. 85	0. 26
123	Whole Brain	210232_at	cell division cycle 42 (GTP-binding protein, 2	Hs. 146409. 1	gb:M35543. 1	CDC42	1	4. 8	197	2. 29	1. 80	0. 10
124	Whole Brain	212559_at	protein kinase, cAMP-dependent, regulatory, ty	Hs. 1519. 0	gb:AU148827_RC			4. 8	366	2. 56	1. 69	0. 18
125	Whole Brain	213965_s_at	KIAA0444 protein	Hs. 158291. 0	gb:AL035406		1	4. 8	168	2. 23	0. 87	0. 28
126	Whole Brain	219005_at	brain-specific membrane-anchored protein	Hs. 5012. 0	gb:NM_012109. 1	BSMAP	19	4. 8	291	2. 46	0. 97	0. 31
127	Whole Brain	219368_at	nucleosome assembly protein 1-like 2	Hs. 66180. 0	gb:NM_021963. 1	NAP1L2	X	4. 8	418	2. 62	1. 47	0. 24
128	Whole Brain	204722_at	voltage-gated sodium channel beta-3 subunit (s	Hs. 4865. 0	gb:AW007335_RC		11	4. 7	570	2. 76	1. 06	0. 36
129	Whole Brain	204966_at	brain-specific angiogenesis inhibitor 2	Hs. 200586. 0	gb:NM_001703. 1	BAI2	1	4. 7	485	2. 69	1. 38	0. 28
130	Whole Brain	205293_x_at	BAI1-associated protein 2	Hs. 7936. 1	gb:AB017120. 1			4. 7	142	2. 15	0. 97	0. 25
131	Whole Brain	205691_at	synaptogyrin 3	Hs. 6467. 0	gb:NM_004209. 2	SYNGR3	16	4. 7	1104	3. 04	1. 03	0. 43
132	Whole Brain	206275_s_at	KIAA0750 gene product	Hs. 314434. 0	gb:NM_014632. 1	KIAA0750	11	4. 7	177	2. 25	1. 15	0. 23
133	Whole Brain	210227_at	discs, large (Drosophila) homolog-associated p	Hs. 113287. 0	gb:AF119817. 1	DLGAP2	8	4. 7	212	2. 33	1. 56	0. 16
134	Whole Brain	213486_at	hypothetical protein DKFZp761N09121	Hs. 6421. 0	gb:BF435376_RC			4. 7	255	2. 41	1. 68	0. 16
135	Whole Brain	214874_at	Homo sapiens mRNA; cDNA DKFZp564G052 (from clo	Hs. 302761. 0	gb:AL050364. 1		2	4. 7	254	2. 40	1. 13	0. 27
136	Whole Brain	219150_s_at	centaurin-alpha	Hs. 135183. 0	gb:NM_006869. 1	GCS1L	7	4. 7	219	2. 34	0. 88	0. 31
137	Whole Brain	219365_s_at	hypothetical protein MGC8407	Hs. 6262. 0	gb:NM_024046. 1	MGC8407	3	4. 7	376	2. 58	1. 78	0. 17
138	Whole Brain	220294_at	neuronal potassium channel alpha subunit	Hs. 13285. 0	gb:NM_014379. 1	KV8. 1	8	4. 7	49	1. 69	0. 22	0. 31
139	Whole Brain	204471_at	growth associated protein 43	Hs. 79000. 0	gb:NM_002045. 1	GAP43	3	4. 6	1037	3. 02	0. 89	0. 46
140	Whole Brain	206039_at	RAB33A, member RAS oncogene family	Hs. 56294. 0	gb:NM_004794. 1	RAB33A	X	4. 6	291	2. 46	1. 55	0. 20
141	Whole Brain	207768_at	early growth response 4	Hs. 3052. 0	gb:NM_001965. 1	EGR4	2	4. 6	137	2. 14	0. 19	0. 43
142	Whole Brain	212772_s_at	KIAA1062 protein	Hs. 94806. 0	gb:AL162060. 1	DKFZp547P1	9	4. 6	296	2. 47	1. 86	0. 13
143	Whole Brain	219219_at	hypothetical protein FLJ20512	Hs. 105606. 0	gb:NM_017854. 1	FLJ20512	19	4. 6	301	2. 48	1. 84	0. 14
144	Whole Brain	219591_at	BM88 antigen	Hs. 22140. 0	gb:NM_016564. 1	LOC51286		4. 6	422	2. 63	1. 77	0. 19
145	Whole Brain	203001_s_at	superiorcervical ganglia, neural specific 10	Hs. 90005. 0	gb:NM_007029. 1	SCGN10	8	4. 5	1157	3. 06	1. 11	0. 43
146	Whole Brain	204247_s_at	cyclin-dependent kinase 5	Hs. 166071. 0	gb:NM_004935. 1	CDK5	7	4. 5	377	2. 58	1. 14	0. 32
147	Whole Brain	204546_at	KIAA0513 gene product	Hs. 301658. 0	gb:NM_014732. 1	KIAA0513	16	4. 5	629	2. 80	1. 67	0. 25
148	Whole Brain	204974_at	RAB3A, member RAS oncogene family	Hs. 27744. 0	gb:AA988241_RC		19	4. 5	487	2. 69	1. 06	0. 36
149	Whole Brain	206051_at	ELAV (embryonic lethal, abnormal vision, Droso	Hs. 75236. 0	gb:NM_021952. 1	ELAVL4		4. 5	553	2. 74	1. 63	0. 24
150	Whole Brain	207659_s_at	myelin-associated oligodendrocyte basic protei	Hs. 169309. 0	gb:NM_006501. 1	MOBP	3	4. 5	409	2. 61	1. 46	0. 25
151	Whole Brain	210408_s_at	copine VI (neuronal)	Hs. 6132. 0	gb:AB009288. 1		14	4. 5	186	2. 27	0. 81	0. 33
152	Whole Brain	212555_at	protein kinase, cAMP-dependent, regulatory, ty	Hs. 1519. 0	gb:AI814660_RC			4. 5	215	2. 33	1. 14	0. 27

Fig. 45

	A	B	C	D	E	F	G	H	I	J	K	L
153	Whole Brain	213849_s_at	protein phosphatase 2 (formerly 2A), regulatory	Hs. 7688. 1	gb:AA974416_RC		1	4. 5	471	2. 67	1. 47	0. 27
154	Whole Brain	214821_at	Homo sapiens clone 23622 mRNA sequence	Hs. 151608. 0	gb:AF052119. 1		4	4. 5	125	2. 10	1. 40	0. 15
155	Whole Brain	205678_at	adaptor-related protein complex 3, beta 2 subur	Hs. 21022. 0	gb:NM_004644. 1	AP3B2	15	4. 4	211	2. 32	1. 42	0. 20
156	Whole Brain	209168_at	glycoprotein M6B	Hs. 5422. 0	gb:AW148844_RC		X	4. 4	383	2. 58	1. 71	0. 20
157	Whole Brain	209618_at	catenin (cadherin-associated protein), delta 2	Hs. 80220. 0	gb:U96136. 1		5	4. 4	384	2. 58	1. 07	0. 34
158	Whole Brain	211836_s_at	myelin oligodendrocyte glycoprotein	Hs. 141308. 1	gb:U18800. 1	MOG	6	4. 4	346	2. 54	1. 21	0. 30
159	Whole Brain	213108_at	calciumcalmodulin-dependent protein kinase (Ca	Hs. 143535. 1	gb:AW131813_RC		5	4. 4	1344	3. 13	1. 52	0. 36
160	Whole Brain	213960_at	Homo sapiens mRNA full length insert cDNA clone	Hs. 13223. 0	gb:T87225_RC		15	4. 4	257	2. 41	1. 69	0. 16
161	Whole Brain	214023_x_at	tubulin, beta polypeptide	Hs. 179661. 4	gb:AL533838_RC		6	4. 4	730	2. 86	1. 07	0. 40
162	Whole Brain	214607_at	p21 (CDKN1A)-activated kinase 3	Hs. 152663. 0	gb:AW085556_RC		X	4. 4	328	2. 52	1. 87	0. 15
163	Whole Brain	200676_s_at	ubiquitin-conjugating enzyme E2L 3	Hs. 108104. 0	gb:NM_003347. 1	UBE2L3	22	4. 3	293	2. 47	1. 95	0. 12
164	Whole Brain	203000_at	superiorcervical ganglia, neural specific 10	Hs. 90005. 0	gb:BF967657_RC		8	4. 3	1123	3. 05	0. 82	0. 52
165	Whole Brain	204953_at	synaptosomal-associated protein, 91 kDa (mouse)	Hs. 12477. 0	gb:NM_014841. 1	KIAA0656	6	4. 3	737	2. 87	1. 08	0. 41
166	Whole Brain	206690_at	amiloride-sensitive cation channel 1, neuronal	Hs. 6517. 0	gb:NM_001094. 2	ACCN1	17	4. 3	121	2. 08	1. 00	0. 25
167	Whole Brain	209372_x_at	tubulin, beta polypeptide	Hs. 179661. 3	gb:BF971587		6	4. 3	423	2. 63	1. 17	0. 34
168	Whole Brain	209883_at	KIAA0584 protein	Hs. 106794. 0	gb:AF288389. 1		1	4. 3	296	2. 47	1. 85	0. 14
169	Whole Brain	221914_at	synapsin I	Hs. 225936. 0	gb:H19843_RC		X	4. 3	766	2. 88	1. 42	0. 34
170	Whole Brain	204684_at	neuronal pentraxin I	Hs. 84154. 0	gb:NM_002522. 1	NPTX1		4. 2	746	2. 87	1. 19	0. 40
171	Whole Brain	205481_at	adenosine A1 receptor	Hs. 77867. 0	gb:NM_000674. 1	ADORA1	1	4. 2	337	2. 53	1. 53	0. 24
172	Whole Brain	205989_s_at	myelin oligodendrocyte glycoprotein	Hs. 141308. 0	gb:NM_002433. 1	MOG	6	4. 2	234	2. 37	0. 85	0. 36
173	Whole Brain	206162_x_at	synaptotagmin V	Hs. 23179. 0	gb:NM_003180. 1	SYT5	18	4. 2	84	1. 92	0. 58	0. 32
174	Whole Brain	209617_s_at	catenin (cadherin-associated protein), delta 2	Hs. 80220. 0	gb:AF035302. 1		5	4. 2	362	2. 56	1. 32	0. 29
175	Whole Brain	209737_at	atrophin-1 interacting protein 1; activin recep	Hs. 22599. 1	gb:AB014605. 1	KIAA0705	7	4. 2	334	2. 52	1. 74	0. 19
176	Whole Brain	213967_at	Homo sapiens clone 24468 mRNA sequence	Hs. 13423. 0	gb:AI634532_RC		8	4. 2	261	2. 42	0. 80	0. 38
177	Whole Brain	217562_at	ESTs, Weakly similar to hypothetical protein H.	Hs. 106642. 0	gb:BF589529_RC		1	4. 2	37	1. 57	0. 27	0. 31
178	Whole Brain	219659_at	ATPase, aminophospholipid transporter-like, Cla	Hs. 130489. 0	gb:AU146927_RC		13	4. 2	233	2. 37	1. 19	0. 28
179	Whole Brain	205113_at	neurofilament 3 (150kD medium)	Hs. 71346. 0	gb:NM_005382. 1	NEF3	8	4. 1	867	2. 94	1. 19	0. 43
180	Whole Brain	206655_s_at	glycoprotein Ib (platelet), beta polypeptide	Hs. 283743. 0	gb:NM_000407. 3	GP1BB	22	4. 1	154	2. 19	0. 75	0. 35
181	Whole Brain	209686_at	S100 calcium-binding protein, beta (neural)	Hs. 83384. 0	gb:BC001766. 1		21	4. 1	440	2. 64	0. 80	0. 45
182	Whole Brain	212308_at	KIAA0627 protein; Drosophila multiple asters (M	Hs. 108614. 0	gb:AL137636. 1	DKFZp434F0	3	4. 1	127	2. 10	1. 35	0. 19
183	Whole Brain	220551_at	differentiation-associated Na-dependent inorgar	Hs. 242821. 0	gb:NM_020346. 1	DNPI	11	4. 1	75	1. 88	1. 11	0. 19
184	Whole Brain	203998_s_at	synaptotagmin I	Hs. 154679. 0	gb:AV723167		12	4	170	2. 23	0. 68	0. 39
185	Whole Brain	204339_s_at	regulator of G-protein signalling 4	Hs. 227571. 0	gb:BC000737. 1		1	4	580	2. 76	1. 50	0. 32
186	Whole Brain	209915_s_at	neurexin 1	Hs. 22998. 0	gb:AB035356. 1		2	4	156	2. 19	0. 90	0. 32
187	Whole Brain	214098_at	KIAA1107 protein	Hs. 21554. 0	gb:AB029030. 1	KIAA1107	1	4	200	2. 30	1. 30	0. 25
188	Whole Brain	215311_at	Homo sapiens mRNA full length insert cDNA clone	Hs. 185701. 0	gb:AL109696. 1		15	4	188	2. 27	1. 52	0. 19
189	Whole Brain	216927_at	KIAA0892 protein	Hs. 112751. 3	gb:AC003030			4	106	2. 03	1. 43	0. 15
190	Whole Brain	219196_at	secretogranin III	Hs. 22215. 0	gb:NM_013243. 1	SCG3	15	4	164	2. 21	0. 38	0. 46

Fig. 46

	A	B	C	D	E	F	G	H	I	J	K	L
191	Whole Brain	221713_s_at			gb:BC006434.1		3	4	272	2.43	1.16	0.32
192	Whole Brain	201340_s_at	ectodermal-neural cortex (with BTB-like domain)	Hs.104925.0	gb:AF010314.1	PIG10	5	3.9	709	2.85	1.43	0.37
193	Whole Brain	202652_at	amyloid beta (A4) precursor protein-binding, fa	Hs.3763.0	gb:NM_001164.1	APBB1	11	3.9	300	2.48	1.75	0.19
194	Whole Brain	204945_at	protein tyrosine phosphatase, receptor type, N	Hs.89655.0	gb:NM_002846.1	PTPRN	2	3.9	194	2.29	1.56	0.18
195	Whole Brain	205651_x_at	cAMP-regulated guanine nucleotide exchange fact	Hs.91971.0	gb:NM_007023.1	CAMP-GEFII	2	3.9	133	2.12	1.10	0.26
196	Whole Brain	209169_at	glycoprotein M6B	Hs.5422.0	gb:N63576_RC		X	3.9	592	2.77	1.31	0.37
197	Whole Brain	209726_at	carbonic anhydrase XI	Hs.22777.0	gb:AB018195.1	ca xi	19	3.9	811	2.91	1.51	0.36
198	Whole Brain	209983_s_at	KIAA0921 protein	Hs.124085.0	gb:AB035266.1	NEUREXIN I	11	3.9	182	2.26	0.71	0.40
199	Whole Brain	213825_at	protein kinase C binding protein 2	Hs.176977.0	gb:AA757419_RC		21	3.9	186	2.27	1.20	0.27
200	Whole Brain	202986_at	KIAA0307 gene product	Hs.6111.0	gb:NM_014862.1	KIAA0307		3.8	492	2.69	1.18	0.40
201	Whole Brain	204729_s_at	syntaxin 1A (brain)	Hs.75671.0	gb:NM_004603.1	STX1A	7	3.8	636	2.80	1.81	0.26
202	Whole Brain	204995_at	cyclin-dependent kinase 5, regulatory subunit 1	Hs.93597.0	gb:AL567411_RC		17	3.8	323	2.51	0.96	0.41
203	Whole Brain	205104_at	syntaphilin	Hs.323833.0	gb:NM_014723.1	KIAA0374	20	3.8	300	2.48	1.32	0.31
204	Whole Brain	205874_at	inositol 1,4,5-trisphosphate 3-kinase A	Hs.2722.0	gb:NM_002220.1	ITPKA	15	3.8	229	2.36	0.84	0.40
205	Whole Brain	211616_s_at			gb:M86841.1	5HT2	13	3.8	87	1.94	1.33	0.16
206	Whole Brain	213268_at	KIAA0833 protein	Hs.26319.0	gb:Z98884		1	3.8	243	2.39	0.86	0.40
207	Whole Brain	201313_at	enolase 2, (gamma, neuronal)	Hs.146580.0	gb:NM_001975.1	ENO2	12	3.7	1498	3.18	1.07	0.57
208	Whole Brain	206984_s_at	Ric (Drosophila)-like, expressed in neurons	Hs.47626.0	gb:NM_002930.1	RIN	18	3.7	153	2.18	0.60	0.42
209	Whole Brain	209469_at	glycoprotein M6A	Hs.75819.0	gb:BF939489_RC		4	3.7	1292	3.11	1.17	0.52
210	Whole Brain	209991_x_at	G protein-coupled receptor 51	Hs.198612.0	gb:AF069755.1	HG20	9	3.7	426	2.63	1.50	0.31
211	Whole Brain	210263_at	potassium voltage-gated channel, subfamily F, m	Hs.23735.0	gb:AF029780.1		2	3.7	92	1.96	0.89	0.29
212	Whole Brain	210292_s_at	protocadherin 11	Hs.159156.2	gb:AF332218.1	PCDH11	X	3.7	98	1.99	1.19	0.21
213	Whole Brain	213045_at	KIAA0561 protein	Hs.173864.0	gb:AB011133.1	KIAA0561	19	3.7	502	2.70	1.61	0.29
214	Whole Brain	214839_at	Homo sapiens clone 23687 mRNA sequence	Hs.12513.0	gb:AF052108.1		8	3.7	154	2.19	1.36	0.22
215	Whole Brain	215241_at	Homo sapiens mRNA for C11ORF25 gene	Hs.91791.0	gb:AJ300461.1			3.7	192	2.28	1.13	0.31
216	Whole Brain	218608_at	putative ATPase	Hs.128866.0	gb:NM_022089.1	HSA9947	1	3.7	229	2.36	0.99	0.37
217	Whole Brain	218868_at	actin-related protein 3-beta	Hs.12887.0	gb:NM_020445.1	ARP3BETA	7	3.7	224	2.35	1.40	0.26
218	Whole Brain	201556_s_at	vesicle-associated membrane protein 2 (synaptob	Hs.194534.0	gb:BC002737.1		17	3.6	387	2.59	1.75	0.24
219	Whole Brain	206826_at	peripheral myelin protein 2	Hs.2868.0	gb:NM_002677.1	PMP2	8	3.6	113	2.05	1.20	0.24
220	Whole Brain	205320_at	adenomatous polyposis coli like	Hs.20912.0	gb:NM_005883.1	APCL	18	3.5	137	2.14	0.85	0.37
221	Whole Brain	205331_s_at	SGC32445 protein	Hs.28088.0	gb:NM_016606.1	LOC51308	5	3.5	98	1.99	1.16	0.24
222	Whole Brain	206456_at	gamma-aminobutyric acid (GABA) A receptor, alph	Hs.24969.0	gb:NM_000810.2	GABRA5	15	3.5	314	2.50	0.75	0.50
223	Whole Brain	209767_s_at	glycoprotein Ib (platelet), beta polypeptide	Hs.283743.1	gb:AA702163_RC		22	3.5	125	2.10	0.87	0.35
224	Whole Brain	209859_at	KIAA0282 protein	Hs.75090.0	gb:AF220036.1	TRIM9	14	3.5	235	2.37	1.09	0.37
225	Whole Brain	210315_at	synapsin II	Hs.6439.0	gb:AF077737.1	SYN2	3	3.5	110	2.04	1.05	0.28
226	Whole Brain	211891_s_at	Rho guanine nucleotide exchange factor (GEF) 4	Hs.6066.1	gb:AB042199.1	Asef		3.5	129	2.11	1.51	0.17
227	Whole Brain	212641_at	human immunodeficiency virus type I enhancer-bi	Hs.75063.0	gb:AL023584		6	3.5	182	2.26	1.52	0.21
228	Whole Brain	213938_at	KIAA0378 protein	Hs.7006.0	gb:Z38645_RC		3	3.5	180	2.26	0.80	0.42

Fig. 47

	A	B	C	D	E	F	G	H	I	J	K	L
229	Whole Brain	219300_s_at	cell recognition molecule Caspr2	Hs. 106552. 0	gb:AB020675. 1	KIAA0868	7	3. 5	143	2. 16	1. 06	0. 32
230	Whole Brain	219738_s_at	protocadherin 9	Hs. 12450. 0	gb:NM_020403. 1	PCDH9	13	3. 5	332	2. 52	1. 02	0. 42
231	Whole Brain	202254_at	KIAA0440 protein	Hs. 172180. 0	gb:AB007900. 1	KIAA0440	14	3. 4	242	2. 38	1. 21	0. 34
232	Whole Brain	203390_s_at	kinesin family member 3C	Hs. 21611. 0	gb:NM_002254. 1	KIF3C	2	3. 4	514	2. 71	1. 74	0. 29
233	Whole Brain	204869_at	proprotein convertase subtilisinkexin type 2	Hs. 93164. 0	gb:AL031664		20	3. 4	237	2. 38	1. 41	0. 29
234	Whole Brain	205638_at	brain-specific angiogenesis inhibitor 3	Hs. 8074. 0	gb:NM_001704. 1	BAI3	6	3. 4	235	2. 37	1. 30	0. 31
235	Whole Brain	206850_at	RAS-related on chromsome 22	Hs. 73088. 0	gb:NM_006477. 1	RRP22	22	3. 4	388	2. 59	1. 66	0. 27
236	Whole Brain	209755_at	KIAA0479 protein	Hs. 158244. 0	gb:AF288395. 1		1	3. 4	467	2. 67	1. 31	0. 40
237	Whole Brain	209914_s_at	neurexin 1	Hs. 22998. 0	gb:AW149405_RC		2	3. 4	117	2. 07	0. 95	0. 33
238	Whole Brain	214432_at	ATPase, Na+K+ transporting, alpha 3 polypeptide	Hs. 274371. 0	gb:NM_000703. 1	ATP1A3	19	3. 4	341	2. 53	1. 11	0. 42
239	Whole Brain	218675_at	organic cation transporter	Hs. 283404. 0	gb:NM_020372. 1	LOC57100	14	3. 4	469	2. 67	1. 36	0. 39
240	Whole Brain	37566_at						3. 3	109	2. 04	0. 96	0. 33
241	Whole Brain	41577_at						3. 3	287	2. 46	1. 10	0. 41
242	Whole Brain	203894_at	tubulin, gamma 2	Hs. 279669. 0	gb:NM_016437. 1	TUBG2	17	3. 3	227	2. 36	1. 68	0. 21
243	Whole Brain	204311_at	ATPase, Na+K+ transporting, beta 2 polypeptide	Hs. 78854. 0	gb:NM_001678. 1	ATP1B2	17	3. 3	244	2. 39	1. 46	0. 29
244	Whole Brain	205029_s_at	fatty acid binding protein 7, brain	Hs. 26770. 0	gb:NM_001446. 1	FABP7	6	3. 3	148	2. 17	0. 50	0. 50
245	Whole Brain	205751_at	SH3-domain GRB2-like 2	Hs. 75149. 0	gb:NM_003026. 1	SH3GL2	9	3. 3	921	2. 96	1. 18	0. 54
246	Whole Brain	207100_s_at	vesicle-associated membrane protein 1 (synaptob	Hs. 20021. 0	gb:NM_016830. 1	VAMP1	12	3. 3	131	2. 12	1. 11	0. 30
247	Whole Brain	207135_at	5-hydroxytryptamine (serotonin) receptor 2A	Hs. 298623. 0	gb:NM_000621. 1	HTR2A	13	3. 3	202	2. 31	1. 58	0. 22
248	Whole Brain	211006_s_at	potassium voltage-gated channel, Shab-related s	Hs. 84244. 0	gb:L02840. 1		20	3. 3	221	2. 34	1. 57	0. 23
249	Whole Brain	213678_at	Homo sapiens germline mRNA sequence	Hs. 12840. 0	gb:F09448_RC		6	3. 3	234	2. 37	1. 61	0. 23
250	Whole Brain	214811_at	KIAA0318 protein	Hs. 65746. 0	gb:AB002316. 1	KIAA0318	12	3. 3	193	2. 29	1. 20	0. 33
251	Whole Brain	215056_at	Human clone 23695 mRNA sequence	Hs. 90798. 0	gb:AI267546_RC		1	3. 3	263	2. 42	1. 70	0. 22
252	Whole Brain	203724_s_at	KIAA0871 protein	Hs. 7972. 0	gb:NM_014961. 1	KIAA0871	4	3. 2	275	2. 44	1. 26	0. 37
253	Whole Brain	204552_at	inositol polyphosphate-4-phosphatase, type I, 1	Hs. 32944. 0	gb:AA355179		2	3. 2	140	2. 15	1. 57	0. 18
254	Whole Brain	205050_s_at	mitogen-activated protein kinase 8 interacting	Hs. 80545. 0	gb:NM_012324. 1	MAPK8IP2	22	3. 2	191	2. 28	1. 38	0. 28
255	Whole Brain	205825_at	proprotein convertase subtilisinkexin type 1	Hs. 78977. 0	gb:NM_000439. 2	PCSK1	5	3. 2	165	2. 22	0. 72	0. 47
256	Whole Brain	212750_at	KIAA0823 protein	Hs. 45719. 0	gb:AB020630. 1	KIAA0823	20	3. 2	489	2. 69	1. 35	0. 42
257	Whole Brain	213436_at	cannabinoid receptor 1 (brain)	Hs. 75110. 1	gb:U73304		6	3. 2	129	2. 11	1. 43	0. 21
258	Whole Brain	214120_at	ret finger protein-like 1 antisense	Hs. 102576. 0	gb:AW157535_RC		22	3. 2	267	2. 43	0. 78	0. 51
259	Whole Brain	214495_at	calcium channel, voltage-dependent, gamma subur	Hs. 268545. 0	gb:NM_006078. 1	CACNG2	22	3. 2	97	1. 99	1. 24	0. 23
260	Whole Brain	214956_at	Homo sapiens clone IMAGE 30714	Hs. 177837. 0	gb:AF090101. 1_RC			3. 2	209	2. 32	1. 25	0. 34
261	Whole Brain	215972_at	Homo sapiens clone 24820 mRNA sequence	Hs. 146312. 0	gb:AF070547. 1		5	3. 2	118	2. 07	1. 17	0. 28
262	Whole Brain	216073_at	Homo sapiens mRNA full length insert cDNA clone	Hs. 21597. 0	gb:AL109678. 1		15	3. 2	74	1. 87	1. 24	0. 20
263	Whole Brain	219714_s_at	calcium channel, voltage-dependent, alpha 2del	Hs. 22958. 0	gb:NM_018398. 1	HSA272268	3	3. 2	130	2. 11	1. 06	0. 33
264	Whole Brain	221082_s_at	hypothetical protein FLJ13556 similar to N-myc	Hs. 240615. 0	gb:NM_022477. 1	FLJ13556	20	3. 2	270	2. 43	1. 84	0. 18
265	Whole Brain	201246_s_at	hypothetical protein FLJ20113	Hs. 108504. 0	gb:NM_017670. 1	FLJ20113	11	3. 1	77	1. 89	0. 95	0. 30
266	Whole Brain	203018_s_at	KIAA0923 protein	Hs. 22587. 0	gb:AU152583_RC		1	3. 1	177	2. 25	1. 26	0. 32

Fig. 48

	A	B	C	D	E	F	G	H	I	J	K	L
267	Whole Brain	203019_x_at	KIAA0923 protein	Hs. 22587. 0	gb:NM_014021. 1	KIAA0923	1	3. 1	178	2. 25	1. 48	0. 24
268	Whole Brain	203798_s_at	visinin-like 1	Hs. 2288. 0	gb:NM_003385. 1	VSNL1	2	3. 1	606	2. 78	1. 03	0. 57
269	Whole Brain	206780_at	glutamate decarboxylase 2 (pancreatic islets ar	Hs. 170808. 0	gb:NM_000818. 1	GAD2	10	3. 1	385	2. 59	1. 50	0. 35
270	Whole Brain	207014_at	gamma-aminobutyric acid (GABA) A receptor, alp	Hs. 91343. 0	gb:NM_000807. 1	GABRA2	4	3. 1	129	2. 11	1. 27	0. 27
271	Whole Brain	207379_at	EGF-like repeats and discoidin I-like domains 3	Hs. 129764. 0	gb:NM_005711. 1	EDIL3	5	3. 1	136	2. 13	1. 35	0. 25
272	Whole Brain	208102_s_at	pleckstrin and Sec7 domain protein	Hs. 154658. 0	gb:NM_002779. 1	PSD		3. 1	102	2. 01	1. 17	0. 27
273	Whole Brain	209631_s_at	G protein-coupled receptor 37 (endothelin recep	Hs. 27747. 0	gb:U87460. 1		9	3. 1	457	2. 66	1. 05	0. 52
274	Whole Brain	209982_s_at	KIAA0921 protein	Hs. 124085. 0	gb:AA608820_RC		11	3. 1	148	2. 17	0. 91	0. 40
275	Whole Brain	212309_at	KIAA0627 protein; Drosophila multiple asters (M	Hs. 108614. 0	gb:AV725315		3	3. 1	251	2. 40	1. 60	0. 26
276	Whole Brain	213609_s_at	seizure related gene 6 (mouse)-like	Hs. 194766. 1	gb:AB023144. 1	KIAA0927	22	3. 1	201	2. 30	1. 48	0. 26
277	Whole Brain	213904_at	Homo sapiens mRNA; cDNA DKFZp547E184 (from clor	Hs. 8077. 0	gb:AL390170. 1		9	3. 1	435	2. 64	0. 93	0. 54
278	Whole Brain	216967_at	growth associated protein 43	Hs. 79000. 1	gb:AF279774. 1_RC		3	3. 1	79	1. 90	1. 15	0. 24
279	Whole Brain	206849_at	gamma-aminobutyric acid (GABA) A receptor, gamm	Hs. 7195. 0	gb:NM_000816. 1	GABRG2	5	3	171	2. 23	1. 59	0. 22
280	Whole Brain	210100_s_at	ATP-binding cassette, sub-family A (ABC1), memb	Hs. 121561. 0	gb:AF327657. 1	ABCA2	9	3	100	2. 00	1. 18	0. 27
281	Whole Brain	219225_at	hypothetical protein FLJ11413	Hs. 12247. 0	gb:NM_024554. 1	FLJ11413	1	3	449	2. 65	1. 50	0. 38
282	Whole Brain	219463_at	similar to S68401 (cattle) glucose induced gene	Hs. 22920. 0	gb:NM_012261. 1	HS1119D91	20	3	439	2. 64	1. 41	0. 41
283	Whole Brain	210871_x_at	KIAA0923 protein	Hs. 22587. 1	gb:AL133046. 1	DKFZp434C0	1	2. 9	201	2. 30	1. 55	0. 27
284	Whole Brain	213439_x_at	RaP2 interacting protein 8	Hs. 6755. 1	gb:AI197870_RC		17	2. 9	305	2. 48	1. 44	0. 36
285	Whole Brain	215728_s_at	cytosolic acyl coenzyme A thioester hydrolase	Hs. 8679. 2	gb:AL031848		1	2. 9	161	2. 21	1. 25	0. 33
286	Whole Brain	205852_at	cyclin-dependent kinase 5, regulatory subunit 2	Hs. 158460. 0	gb:R51311_RC		2	2. 8	94	1. 97	0. 82	0. 41
287	Whole Brain	210383_at	sodium channel, voltage-gated, type I, alpha pc	Hs. 22654. 0	gb:AF225985. 1	SCN1A	2	2. 8	65	1. 81	0. 66	0. 40
288	Whole Brain	204375_at	KIAA0726 gene product	Hs. 107809. 0	gb:NM_014718. 1	KIAA0726		2. 7	172	2. 24	1. 48	0. 28
289	Whole Brain	214217_at	Human clone 23560 mRNA sequence	Hs. 79981. 0	gb:D60132_RC		11	2. 7	105	2. 02	1. 16	0. 33
290	Whole Brain	209867_s_at	KIAA0768 protein	Hs. 21917. 0	gb:AF307080. 1	LEC3	4	2. 6	114	2. 06	1. 08	0. 38
291	Whole Brain	213745_at	KIAA0534 protein	Hs. 196012. 0	gb:AW151108_RC		10	2. 5	108	2. 03	1. 17	0. 35
292	Whole Brain	214998_at	Homo sapiens clone IMAGE 23915	Hs. 170241. 0	gb:AF090100. 1_RC			2. 5	99	1. 99	1. 24	0. 30
293	Whole Brain	203839_s_at	activated p21cdc42Hs kinase	Hs. 153937. 0	gb:NM_005781. 2	ACK1	3	2. 4	168	2. 23	1. 42	0. 34
294	Whole Brain	210721_s_at	KIAA1264 protein	Hs. 32539. 0	gb:AB040812. 1	PAK5	20	2. 2	61	1. 78	0. 62	0. 54
295	Brain Amygda	207010_at	gamma-aminobutyric acid (GABA) A receptor, beta	Hs. 89768. 0	gb:NM_000812. 2	GABRB1	4	3. 7	307	2. 49	1. 49	0. 27
296	Brain caudat	220313_at	G-protein coupled receptor 88	Hs. 170053. 0	gb:NM_022049. 1	GPR88	1	6. 2	1481	3. 17	1. 31	0. 30
297	Brain caudat	205501_at	phosphodiesterase 10A	Hs. 128291. 0	gb:AI143879_RC		6	5. 7	984	2. 99	1. 43	0. 27
298	Brain caudat	205362_s_at	prefoldin 4	Hs. 91161. 0	gb:NM_002623. 2	PFDN4	20	4. 1	99	2. 00	1. 43	0. 14
299	Brain caudat	206518_s_at	regulator of G-protein signalling 9	Hs. 117149. 0	gb:NM_003835. 1	RGS9	17	3. 8	193	2. 29	1. 18	0. 29
300	Brain caudat	214655_at	G protein-coupled receptor 6	Hs. 46332. 0	gb:U18549			3. 3	535	2. 73	1. 10	0. 49
301	Brain caudat	38290_at						3. 1	326	2. 51	1. 56	0. 31
302	Brain caudat	207307_at	5-hydroxytryptamine (serotonin) receptor 2C	Hs. 46362. 0	gb:NM_000868. 1	HTR2C	X	3. 1	471	2. 67	0. 95	0. 56
303	Brain caudat	214652_at	dopamine receptor D1	Hs. 2624. 0	gb:X58987. 1		5	2. 5	69	1. 84	0. 75	0. 44
304	Brain corpus	203552_at	mitogen-activated protein kinase kinase kinase	Hs. 246970. 0	gb:AW298170_RC		14	3. 3	259	2. 41	1. 71	0. 21

Fig. 49

	A	B	C	D	E	F	G	H	I	J	K	L
305	Brain corpus	201295_s_at	DKFZP564A122 protein	Hs. 187991. 0	gb:BF111821_RC		17	3. 1	164	2. 22	1. 42	0. 26
306	Brain hippoc	215972_at	Homo sapiens clone 24820 mRNA sequence	Hs. 146312. 0	gb:AF070547. 1		5	3. 1	99	2. 00	1. 16	0. 27
307	Brain cerebe	207933_at	zona pellucida glycoprotein 2 (sperm receptor)	Hs. 73982. 0	gb:NM_003460. 1	ZP2	16	11. 5	184	2. 27	0. 44	0. 16
308	Brain cerebe	213171_s_at	matrix metalloproteinase 24 (membrane-inserted)	Hs. 3743. 2	gb:AL121753		20	7. 3	176	2. 25	0. 94	0. 18
309	Brain cerebe	208457_at	gamma-aminobutyric acid (GABA) A receptor, del	Hs. 113882. 0	gb:NM_000815. 1	GABRD	1	6. 2	759	2. 88	0. 90	0. 32
310	Brain cerebe	64488_at						5. 6	260	2. 41	1. 72	0. 13
311	Brain cerebe	206328_at	cadherin 15, M-cadherin (myotubule)	Hs. 148090. 0	gb:NM_004933. 1	CDH15	NA	5. 6	70	1. 84	0. 48	0. 24
312	Brain cerebe	210400_at	glutamate receptor, ionotropic, N-methyl D-aspa	Hs. 36451. 0	gb:L76224. 1		17	5. 4	423	2. 63	0. 67	0. 36
313	Brain cerebe	214951_at	UDP-N-acetyl-alpha-D-galactosamine: (N-acetylneu	Hs. 159481. 1	gb:AL050358. 1	DKFZp564M1	12	5. 3	159	2. 20	1. 59	0. 12
314	Brain cerebe	206275_s_at	KIAA0750 gene product	Hs. 314434. 0	gb:NM_014632. 1	KIAA0750	11	4. 9	429	2. 63	1. 22	0. 29
315	Brain cerebe	208153_s_at			gb:NM_001447. 1	FAT2	5	4. 8	915	2. 96	1. 39	0. 33
316	Brain cerebe	210341_at	hypothetical protein FLJ10883	Hs. 184628. 1	gb:AB020642. 1	KIAA0835	20	4. 8	275	2. 44	0. 73	0. 36
317	Brain cerebe	220584_at	hypothetical protein FLJ22184	Hs. 288540. 0	gb:NM_025094. 1	FLJ22184	19	4. 8	353	2. 55	0. 91	0. 34
318	Brain cerebe	207060_at	engrailed homolog 2	Hs. 134989. 0	gb:NM_001427. 2	EN2	7	4. 7	328	2. 52	1. 15	0. 29
319	Brain cerebe	215233_at	phosphatidylserine receptor	Hs. 72660. 1	gb:AA351360_RC		17	4. 6	209	2. 32	0. 84	0. 32
320	Brain cerebe	201320_at	SWISNF related, matrix associated, actin depend	Hs. 236030. 0	gb:BF663402		12	4. 5	331	2. 52	1. 93	0. 13
321	Brain cerebe	206163_at	mab-21 (C. elegans)-like 1	Hs. 239506. 0	gb:NM_005584. 1	MAB21L1	13	4. 3	335	2. 52	1. 30	0. 29
322	Brain cerebe	205747_at	cerebellin 1 precursor	Hs. 662. 0	gb:NM_004352. 1	CBLN1	16	4. 2	854	2. 93	1. 32	0. 38
323	Brain cerebe	207182_at	gamma-aminobutyric acid (GABA) A receptor, alp	Hs. 90791. 0	gb:NM_000811. 1	GABRA6		4. 2	192	2. 28	0. 90	0. 33
324	Brain cerebe	216452_at	KIAA1616 protein	Hs. 288911. 0	gb:AB046836. 1	KIAA1616		4. 2	351	2. 55	1. 70	0. 20
325	Brain cerebe	221921_s_at	hypothetical protein FLJ10698	Hs. 6164. 0	gb:AI951798_RC		1	4. 2	242	2. 38	0. 81	0. 37
326	Brain cerebe	206338_at	ELAV (embryonic lethal, abnormal vision, Drosop	Hs. 1701. 0	gb:NM_001420. 2	ELAVL3	19	4. 1	52	1. 71	0. 70	0. 25
327	Brain cerebe	211677_x_at			gb:AF062733. 2	NECL1	1	4. 1	424	2. 63	1. 50	0. 27
328	Brain cerebe	215104_at	Homo sapiens mRNA; cDNA DKFZp761G1913 (from cl	Hs. 245880. 0	gb:AL136557. 1	DKFZp761G1	12	4. 1	185	2. 27	1. 10	0. 28
329	Brain cerebe	221838_at	Homo sapiens cDNA FLJ14307 fis, clone PLACE3000	Hs. 181341. 1	gb:N38751_RC		22	4. 1	125	2. 10	1. 50	0. 14
330	Brain cerebe	205663_at	poly(rC)-binding protein 3	Hs. 121241. 0	gb:NM_020528. 1	PCBP3	21	3. 9	310	2. 49	1. 51	0. 25
331	Brain cerebe	206914_at	class-I MHC-restricted T cell associated molecu	Hs. 159523. 0	gb:NM_019604. 1	CRTAM	11	3. 9	354	2. 55	0. 77	0. 46
332	Brain cerebe	205889_s_at	KIAA0555 gene product	Hs. 43107. 0	gb:NM_014790. 1	KIAA0555	5	3. 8	84	1. 92	1. 34	0. 15
333	Brain cerebe	206230_at	LIM homeobox protein 1	Hs. 157449. 0	gb:NM_005568. 1	LHX1	17	3. 8	56	1. 75	0. 43	0. 35
334	Brain cerebe	215181_at	hypothetical protein similar to cadherin 22	Hs. 264157. 0	gb:AF035300. 1		20	3. 8	462	2. 66	1. 27	0. 37
335	Brain cerebe	214978_s_at	Homo sapiens cDNA FLJ13303 fis, clone OVARC1001	Hs. 300704. 0	gb:AK023365. 1		1	3. 7	548	2. 74	0. 81	0. 52
336	Brain cerebe	213155_at	KIAA0523 protein	Hs. 16032. 0	gb:AB011095. 1	KIAA0523	17	3. 6	342	2. 53	1. 11	0. 39
337	Brain cerebe	206282_at	neurogenic differentiation 1	Hs. 72981. 0	gb:NM_002500. 1	NEUROD1	2	3. 5	561	2. 75	1. 00	0. 50
338	Brain cerebe	213230_at	paraneoplastic antigen	Hs. 78358. 0	gb:AI422335_RC		17	3. 5	388	2. 59	1. 15	0. 41
339	Brain cerebe	214495_at	calcium channel, voltage-dependent, gamma subur	Hs. 268545. 0	gb:NM_006078. 1	CACNG2	22	3. 5	147	2. 17	1. 28	0. 25
340	Brain cerebe	214781_at	plexin A1	Hs. 322844. 1	gb:AL162013. 1	DKFZp761P1	3	3. 5	60	1. 78	0. 31	0. 42
341	Brain cerebe	205646_s_at	paired box gene 6 (aniridia, keratitis)	Hs. 89506. 0	gb:NM_000280. 1	PAX6	11	3. 4	631	2. 80	1. 15	0. 49
342	Brain cerebe	205245_at	Tax interaction protein 40	Hs. 112933. 0	gb:NM_016948. 1	TAX40	16	3. 3	177	2. 25	1. 65	0. 18

Fig. 50

	A	B	C	D	E	F	G	H	I	J	K	L
343	Brain cerebe	212564_at	KIAA0176 protein	Hs. 4935. 0	gb:AA523921_RC			3. 3	503	2. 70	1. 83	0. 26
344	Brain cerebe	214993_at	Homo sapiens clone 24488 mRNA sequence	Hs. 323815. 0	gb:AF070642. 1			3. 3	104	2. 02	0. 75	0. 38
345	Brain cerebe	222352_at	ESTs	Hs. 227170. 0	gb:BE465371_RC		16	3. 3	164	2. 21	1. 51	0. 21
346	Brain cerebe	205795_at	neurexin 3	Hs. 22269. 0	gb:NM_004796. 1	NRXN3	14	3. 2	389	2. 59	0. 83	0. 56
347	Brain cerebe	207831_x_at	deoxyhypusine synthase	Hs. 79064. 0	gb:NM_013407. 1	DHPS	19	3. 2	247	2. 39	1. 32	0. 34
348	Brain cerebe	214903_at	Homo sapiens clone 24527 mRNA sequence	Hs. 25422. 0	gb:AF070580. 1		1	3. 2	519	2. 72	1. 64	0. 34
349	Brain cerebe	221496_s_at	transducer of ERBB2, 2	Hs. 4994. 0	gb:D64109. 1		22	3. 2	387	2. 59	1. 50	0. 34
350	Brain cerebe	213329_at	KIAA0456 protein	Hs. 5003. 0	gb:AA742261_RC		1	3. 1	384	2. 58	1. 51	0. 34
351	Brain cerebe	214597_at	somatostatin receptor 2	Hs. 184841. 0	gb:BC000256. 1		17	3. 1	69	1. 84	1. 13	0. 23
352	Brain cerebe	219114_at	g20 protein	Hs. 21050. 0	gb:NM_016210. 1	LOC51161	3	3	201	2. 30	1. 24	0. 35
353	Brain cerebe	207197_at	Zic family member 3 (odd-paired Drosophila hom	Hs. 111227. 0	gb:NM_003413. 1	ZIC3	X	2. 8	92	1. 96	1. 09	0. 32
354	Brain thalam	206281_at	adenylate cyclase activating polypeptide 1 (pit	Hs. 68137. 0	gb:NM_001117. 2	ADCYAP1		4. 9	220	2. 34	1. 80	0. 11
355	Brain thalam	209324_s_at	regulator of G-protein signalling 16	Hs. 183601. 0	gb:BF304996		1	4. 8	762	2. 88	1. 76	0. 23
356	Brain thalam	210135_s_at	short stature homeobox 2	Hs. 55967. 2	gb:AF022654. 1	OG12		3. 9	248	2. 40	1. 12	0. 32
357	Brain thalam	212759_s_at	transcription factor 7-like 2 (T-cell specific,	Hs. 173638. 0	gb:AI703074_RC		10	3. 8	236	2. 37	0. 91	0. 38
358	Brain thalam	216073_at	Homo sapiens mRNA full length insert cDNA clone	Hs. 21597. 0	gb:AL109678. 1		15	3. 8	194	2. 29	1. 32	0. 25
359	Pituitary Gl	205720_at	proopiomelanocortin (adrenocorticotropin beta-l	Hs. 1897. 0	gb:NM_000939. 1	POMC	2	17. 2	6786	3. 83	0. 67	0. 18
360	Pituitary Gl	214529_at	thyroid stimulating hormone, beta	Hs. 53213. 0	gb:NM_000549. 1	TSHB	1	13. 5	2486	3. 40	1. 86	0. 11
361	Pituitary Gl	214471_x_at	luteinizing hormone beta polypeptide	Hs. 154704. 0	gb:NM_000894. 1	LHB	19	7. 2	2499	3. 40	1. 29	0. 29
362	Pituitary Gl	207825_s_at	growth hormone releasing hormone receptor	Hs. 767. 0	gb:NM_000823. 1	GHRHR	7	6. 5	1085	3. 04	1. 02	0. 31
363	Pituitary Gl	206838_at	T-box 19	Hs. 50403. 0	gb:NM_005149. 1	TBX19	1	6. 4	645	2. 81	1. 03	0. 28
364	Pituitary Gl	220288_at	myosin XV	Hs. 127561. 0	gb:NM_016239. 1	MYO15	17	5. 5	362	2. 56	1. 69	0. 16
365	Pituitary Gl	207261_at	cyclic nucleotide gated channel alpha 3	Hs. 234785. 0	gb:NM_001298. 1	CNGA3	2	5. 1	95	1. 98	0. 52	0. 29
366	Pituitary Gl	207695_s_at	immunoglobulin superfamily, member 1	Hs. 22111. 0	gb:NM_001555. 1	IGSF1		5. 1	1593	3. 20	1. 48	0. 34
367	Pituitary Gl	214489_at	follicle stimulating hormone, beta polypeptide	Hs. 36975. 0	gb:NM_000510. 1	FSHB	11	4. 9	513	2. 71	1. 27	0. 30
368	Pituitary Gl	211544_s_at	growth hormone releasing hormone receptor	Hs. 767. 5	gb:AB058895. 1	GHRHRpsv	7	4. 5	646	2. 81	1. 51	0. 29
369	Pituitary Gl	205387_s_at	chorionic gonadotropin, beta polypeptide	Hs. 172944. 0	gb:NM_000737. 1	CGB	19	4. 2	127	2. 10	0. 48	0. 39
370	Pituitary Gl	211522_s_at	gonadotropin-releasing hormone receptor	Hs. 73064. 0	gb:L03380. 1		4	4. 2	92	1. 96	0. 50	0. 35
371	Pituitary Gl	219848_s_at	KIAA0798 gene product	Hs. 178471. 0	gb:NM_014650. 1	KIAA0798	19	4. 1	235	2. 37	1. 76	0. 15
372	Pituitary Gl	208502_s_at	paired-like homeodomain transcription factor 1	Hs. 84136. 1	gb:NM_002653. 1	PITX1	5	3. 8	617	2. 79	0. 66	0. 57
373	Pituitary Gl	211545_at	growth hormone releasing hormone receptor	Hs. 767. 5	gb:AB058895. 1	GHRHRpsv	7	3. 8	197	2. 29	0. 90	0. 37
374	Pituitary Gl	219355_at	hypothetical protein FLJ10178	Hs. 274267. 0	gb:NM_018015. 2	FLJ10178	X	3. 7	290	2. 46	0. 76	0. 46
375	Pituitary Gl	207624_s_at	retinitis pigmentosa GTPase regulator	Hs. 153614. 0	gb:NM_000328. 1	RPGR		3. 6	122	2. 09	1. 49	0. 16
376	Pituitary Gl	206841_at	phosphodiesterase 6H, cGMP-specific, cone, gam	Hs. 54471. 0	gb:NM_006205. 1	PDE6H		3. 5	106	2. 03	0. 73	0. 37
377	Pituitary Gl	222095_s_at	KIAA0700 protein	Hs. 13999. 1	gb:AW450345_RC		1	3. 5	155	2. 19	0. 86	0. 38
378	Pituitary Gl	207846_at	POU domain, class 1, transcription factor 1 (Pi	Hs. 89394. 0	gb:NM_000306. 1	POU1F1	3	3. 2	122	2. 09	0. 55	0. 48
379	Pituitary Gl	209587_at	paired-like homeodomain transcription factor 1	Hs. 84136. 0	gb:U70370. 1	Bft	5	3. 2	192	2. 28	0. 83	0. 45
380	Pituitary Gl	214601_at	tryptophan hydroxylase (tryptophan 5-monooxyger	Hs. 144563. 0	gb:AI350339_RC		11	3. 2	165	2. 22	0. 40	0. 57

Fig. 51

	A	B	C	D	E	F	G	H	I	J	K	L
381	Pituitary Gl	220484_at	hypothetical protein FLJ11006	Hs. 49344. 0	gb:NM_018298. 1	FLJ11006	1	3. 2	173	2. 24	1. 09	0. 36
382	Pituitary Gl	222293_at	ESTs, Highly similar to F22162 1 H. sapiens	Hs. 312723. 0	gb:AW204383_RC		19	2. 4	76	1. 88	1. 05	0. 35
383	Pituitary Gl	203050_at	tumor protein p53-binding protein, 1	Hs. 170263. 0	gb:NM_005657. 1	TP53BP1	15	2	398	2. 60	1. 97	0. 32
384	Spinal Cord	210280_at	myelin protein zero (Charcot-Marie-Tooth neuro	Hs. 93883. 0	gb:D10537. 1			6. 2	851	2. 93	1. 55	0. 22
385	Spinal Cord	204764_at	farnesyltransferase, CAAX box, beta	Hs. 322659. 0	gb:NM_002028. 1	FNTB	14	4. 1	278	2. 44	1. 91	0. 13
386	Spinal Cord	1773_at						3. 9	231	2. 36	1. 68	0. 18
387	Spinal Cord	219895_at	hypothetical protein FLJ20716	Hs. 144517. 0	gb:NM_017938. 1	FLJ20716	X	3. 3	168	2. 22	1. 33	0. 28
388	Salivary Gla	208555_x_at	cystatin SA	Hs. 247955. 0	gb:NM_001322. 1	CST2		10	12255	4. 09	1. 12	0. 30
389	Salivary Gla	206224_at	cystatin SN	Hs. 123114. 0	gb:NM_001898. 1	CST1	20	8. 5	13967	4. 15	0. 96	0. 38
390	Salivary Gla	207925_at	cystatin D	Hs. 121489. 0	gb:NM_001900. 1	CST5		6. 1	9141	3. 96	1. 16	0. 46
391	Salivary Gla	216359_at	Homo sapiens (clone MG2-5-12) mucin (MG2) mRNA,	Hs. 103944. 0	gb:L13283. 1_RC		4	4. 8	201	2. 30	1. 36	0. 20
392	Salivary Gla	206827_s_at	Alu-binding protein with zinc finger domain	Hs. 289104. 0	gb:NM_014274. 1	ABPZF		4. 7	177	2. 25	0. 58	0. 35
393	Salivary Gla	220380_at	deoxyribonuclease II beta	Hs. 129142. 0	gb:NM_021233. 1	LOC58511	1	4. 2	340	2. 53	1. 07	0. 35
394	Salivary Gla	216994_s_at	runt-related transcription factor 2	Hs. 121895. 1	gb:L40992. 1	CBFA1	6	4	87	1. 94	1. 38	0. 14
395	Salivary Gla	217999_s_at	pleckstrin homology-like domain, family A, mem	Hs. 82101. 0	gb:NM_007350. 1	PHLDA1	12	3. 6	205	2. 31	1. 59	0. 20
396	Salivary Gla	212246_at	Human transposon-like element mRNA	Hs. 84775. 0	gb:BE880828		2	3. 5	394	2. 60	1. 74	0. 25
397	Salivary Gla	202700_s_at	KIAA0792 gene product	Hs. 119387. 0	gb:NM_014698. 1	KIAA0792	1	3. 1	132	2. 12	1. 39	0. 24
398	Salivary Gla	207380_x_at	NADPH oxidase 1	Hs. 132370. 2	gb:NM_013954. 1	NOX1	X	3. 1	190	2. 28	1. 61	0. 22
399	Salivary Gla	206896_s_at	guanine nucleotide binding protein (G protein),	Hs. 127828. 0	gb:NM_005145. 1	GNG7	NA	3	167	2. 22	1. 39	0. 28
400	Salivary Gla	206977_at	parathyroid hormone	Hs. 37045. 0	gb:NM_000315. 1	PTH	11	3	65	1. 81	0. 53	0. 43
401	Salivary Gla	207591_s_at	SWISNF related, matrix associated, actin depen	Hs. 123090. 0	gb:NM_006015. 1	SMARCF1		2. 9	94	1. 97	1. 05	0. 32
402	Thymus	210487_at	deoxynucleotidyltransferase, terminal	Hs. 272537. 0	gb:M11722. 1	LA0024A	10	12. 1	442	2. 64	0. 74	0. 16
403	Thymus	206804_at	CD3G antigen, gamma polypeptide (TiT3 complex)	Hs. 2259. 0	gb:NM_000073. 1	CD3G	11	10. 1	232	2. 37	0. 93	0. 14
404	Thymus	207445_s_at	chemokine (C-C motif) receptor 9	Hs. 225946. 0	gb:AF145439. 1	CCR9	3	8. 5	772	2. 89	1. 49	0. 16
405	Thymus	206181_at	signaling lymphocytic activation molecule	Hs. 32970. 0	gb:NM_003037. 1	SLAM	1	6. 6	577	2. 76	1. 67	0. 17
406	Thymus	206591_at	recombination activating gene 1	Hs. 73958. 0	gb:NM_000448. 1	RAG1	11	6. 4	673	2. 83	1. 02	0. 28
407	Thymus	215731_s_at	M-phase phosphoprotein 9	Hs. 86178. 2	gb:X98258. 1	mpp9	12	6. 4	59	1. 77	0. 42	0. 21
408	Thymus	208592_s_at			gb:NM_030893. 1	CD1E	1	6. 2	307	2. 49	1. 28	0. 20
409	Thymus	206749_at	CD1B antigen, b polypeptide	Hs. 1310. 0	gb:NM_001764. 1	CD1B	1	5. 6	788	2. 90	1. 16	0. 31
410	Thymus	209574_s_at	chromosome 18 open reading frame 1	Hs. 153498. 2	gb:AI349506_RC		18	5. 4	281	2. 45	1. 71	0. 14
411	Thymus	210116_at	SH2 domain protein 1A, Duncans disease (lympho	Hs. 151544. 0	gb:AF072930. 1		X	5. 3	410	2. 61	0. 69	0. 36
412	Thymus	210556_at	nuclear factor of activated T-cells, cytoplasmi	Hs. 172674. 2	gb:U85430. 1		16	5	255	2. 41	1. 47	0. 19
413	Thymus	221965_at	M-phase phosphoprotein 9	Hs. 86178. 1	gb:AI990326_RC		12	5	268	2. 43	1. 95	0. 10
414	Thymus	205254_x_at	transcription factor 7 (T-cell specific, HMG-b	Hs. 169294. 0	gb:AW027359_RC		5	4. 8	177	2. 25	1. 05	0. 25
415	Thymus	215784_at	CD1E antigen, e polypeptide	Hs. 249217. 0	gb:AA309511		1	4. 8	1219	3. 09	0. 90	0. 45
416	Thymus	207979_s_at	CD8 antigen, beta polypeptide 1 (p37)	Hs. 2299. 0	gb:NM_004931. 1	CD8B1		4. 3	648	2. 81	1. 66	0. 27
417	Thymus	210001_s_at	JAK binding protein	Hs. 50640. 0	gb:AB005043. 1		16	4. 2	196	2. 29	1. 55	0. 18
418	Thymus	211210_x_at	SH2 domain protein 1A, Duncans disease (lympho	Hs. 151544. 2	gb:AF100539. 1	SH2D1A	X	4	305	2. 48	1. 53	0. 24

Fig. 52

	A	B	C	D	E	F	G	H	I	J	K	L
419	Thymus	216757_at	Homo sapiens cDNA: FLJ21342 fis, clone COL02673	Hs. 306739. 0	gb:AK024995. 1_RC			4	237	2. 38	1. 77	0. 15
420	Thymus	204529_s_at	KIAA0808 gene product	Hs. 184297. 0	gb:AI961231_RC		8	3. 8	283	2. 45	1. 59	0. 22
421	Thymus	205987_at	CD1C antigen, c polypeptide	Hs. 1311. 0	gb:NM_001765. 1	CD1C	1	3. 8	322	2. 51	1. 07	0. 38
422	Thymus	208165_s_at	protease, serine, 16 (thymus)	Hs. 274407. 0	gb:NM_005865. 1	PRSS16	6	3. 8	655	2. 82	1. 58	0. 32
423	Thymus	211005_at	linker for activation of T cells	Hs. 83496. 1	gb:AF036906. 1	LAT		3. 8	351	2. 55	1. 65	0. 23
424	Thymus	215332_s_at	CD8 antigen, beta polypeptide 1 (p37)	Hs. 2299. 4	gb:AW296309_RC			3. 8	124	2. 09	0. 68	0. 37
425	Thymus	211272_s_at	diacylglycerol kinase, alpha (80kD)	Hs. 172690. 1	gb:AF064771. 1	DAGK1	12	3. 7	163	2. 21	1. 49	0. 19
426	Thymus	204890_s_at	lymphocyte-specific protein tyrosine kinase	Hs. 1765. 0	gb:U07236. 1	LCK	1	3. 4	435	2. 64	1. 29	0. 39
427	Thymus	211211_x_at	SH2 domain protein 1A, Duncans disease (lymphop	Hs. 151544. 3	gb:AF100542. 1	SH2D1A	X	3. 4	270	2. 43	1. 53	0. 27
428	Thymus	221897_at	hypothetical protein FLJ23053	Hs. 94037. 1	gb:AA205660_RC		5	3. 4	162	2. 21	1. 63	0. 17
429	Thymus	217147_s_at	T-cell receptor interacting molecule	Hs. 138701. 1	gb:AJ240085. 1	TRIM	3	3. 3	212	2. 33	0. 74	0. 48
430	Thymus	206551_x_at	hypothetical protein FLJ20059	Hs. 246875. 0	gb:NM_017644. 1	FLJ20059	3	3. 2	204	2. 31	1. 66	0. 20
431	Thymus	207996_s_at	chromosome 18 open reading frame 1	Hs. 153498. 0	gb:NM_004338. 1	C18ORF1	18	3. 2	241	2. 38	1. 54	0. 26
432	Thymus	217143_s_at	T cell receptor delta locus	Hs. 2014. 2	gb:X06557. 1		14	3. 1	492	2. 69	1. 45	0. 41
433	Thymus	215117_at	recombination activating gene 2	Hs. 159376. 0	gb:AW058148_RC		11	3	83	1. 92	0. 66	0. 41
434	Thymus	219918_s_at	hypothetical protein FLJ10517	Hs. 279797. 0	gb:NM_018123. 1	FLJ10517	1	3	258	2. 41	0. 72	0. 56
435	Thymus	215907_at	Homo sapiens cDNA: FLJ23540 fis, clone LNG0823	Hs. 293931. 0	gb:AK027193. 1		6	2. 9	177	2. 25	1. 24	0. 35
436	Thymus	210948_s_at	lymphoid enhancer binding factor-1	Hs. 44865. 0	gb:AF294627. 1	LEF1	4	2. 8	164	2. 21	0. 92	0. 46
437	Thymus	222145_at	Homo sapiens cDNA: FLJ23572 fis, clone LNG1240	Hs. 201593. 0	gb:AK027225. 1		16	2. 2	161	2. 21	1. 17	0. 48
438	Thyroid	206529_x_at	solute carrier family, member 4	Hs. 159275. 0	gb:NM_000441. 1	SLC26A4	7	6. 7	520	2. 72	1. 15	0. 23
439	Thyroid	210342_s_at	thyroid peroxidase	Hs. 2041. 0	gb:M17755. 2	TPO	2	6. 6	3422	3. 53	0. 80	0. 41
440	Thyroid	214977_at	Homo sapiens cDNA FLJ13790 fis, clone THYRO100	Hs. 301407. 0	gb:AK023852. 1		8	6. 6	277	2. 44	0. 54	0. 29
441	Thyroid	215442_s_at	thyroid stimulating hormone receptor	Hs. 123078. 1	gb:BE740743		14	6. 2	198	2. 30	0. 34	0. 32
442	Thyroid	210055_at	thyroid stimulating hormone receptor	Hs. 123078. 0	gb:BE045816_RC		14	5. 3	652	2. 81	0. 68	0. 40
443	Thyroid	211122_s_at	small inducible cytokine subfamily B (Cys-X-Cys	Hs. 103982. 1	gb:AF002985. 1	H174		4. 6	175	2. 24	1. 22	0. 22
444	Thyroid	215443_at	thyroid stimulating hormone receptor	Hs. 123078. 1	gb:BE740743		14	4. 5	257	2. 41	0. 45	0. 43
445	Thyroid	203699_s_at	deiodinase, iodothyronine, type II	Hs. 154424. 0	gb:U53506. 1		14	4. 4	396	2. 60	1. 12	0. 34
446	Thyroid	204933_s_at	tumor necrosis factor receptor superfamily, mem	Hs. 81791. 0	gb:NM_002546. 1	TNFRSF11B	8	4. 1	207	2. 32	0. 54	0. 44
447	Thyroid	210673_x_at	thyroid transcription factor 1	Hs. 197764. 3	gb:D50740. 1		14	4	129	2. 11	0. 68	0. 36
448	Thyroid	213651_at	phosphatidylinositol (4, 5) bisphosphate 5-phosp	Hs. 21492. 0	gb:AI935720_RC		22	4	352	2. 55	1. 20	0. 34
449	Thyroid	204932_at	tumor necrosis factor receptor superfamily, mem	Hs. 81791. 0	gb:BF433902_RC		8	3. 9	180	2. 26	1. 09	0. 30
450	Thyroid	205139_s_at	uronyl 2-sulfotransferase	Hs. 134015. 0	gb:NM_005715. 1	UST	6	3. 7	251	2. 40	1. 86	0. 15
451	Thyroid	206912_at	forkhead box E1 (thyroid transcription factor 2	Hs. 159234. 0	gb:NM_004473. 2	FOXE1	9	3. 7	357	2. 55	0. 96	0. 43
452	Thyroid	203700_s_at	deiodinase, iodothyronine, type II	Hs. 154424. 0	gb:NM_013989. 1	DIO2	14	3. 6	419	2. 62	1. 84	0. 22
453	Thyroid	209292_at	inhibitor of DNA binding 4, dominant negative h	Hs. 34853. 0	gb:AL022726		6	3. 3	533	2. 73	1. 53	0. 36
454	Thyroid	219874_at	hypothetical protein FLJ23188	Hs. 36793. 0	gb:NM_024628. 1	FLJ23188	3	3. 3	156	2. 19	1. 38	0. 25
455	Thyroid	210163_at	small inducible cytokine subfamily B (Cys-X-Cys	Hs. 103982. 0	gb:AF030514. 1			3. 2	168	2. 23	0. 86	0. 43
456	Thyroid	207447_s_at	UDP-N-acetylglucosamine:a-1, 3-D-mannoside beta-	Hs. 227473. 0	gb:NM_013244. 1	HGNT-IV-H		3. 1	77	1. 89	0. 95	0. 30

Fig. 53

	A	B	C	D	E	F	G	H	I	J	K	L
457	Thyroid	216541_x_at	Homo sapiens partial IGVH1 gene for immunoglobu	Hs. 272359. 0	gb:AJ275397			2. 4	226	2. 35	1. 14	0. 50
458	Trachea	206869_at	chondroadherin	Hs. 97220. 0	gb:NM_001267. 1	CHAD	17	9. 6	292	2. 47	0. 71	0. 18
459	Trachea	219837_s_at	cytokine-like protein C17	Hs. 13872. 0	gb:NM_018659. 1	C17		6. 6	2134	3. 33	0. 96	0. 36
460	Trachea	207367_at	ATPase, H+K+ transporting, nongastric, alpha pc	Hs. 1165. 0	gb:NM_001676. 2	ATP12A	13	5. 9	649	2. 81	1. 64	0. 20
461	Trachea	207692_s_at	aggrecan 1 (chondroitin sulfate proteoglycan 1,	Hs. 2159. 1	gb:NM_001135. 1	AGC1	15	4. 8	208	2. 32	1. 68	0. 13
462	Trachea	217404_s_at	collagen, type II, alpha 1 (primary osteoarthri	Hs. 81343. 1	gb:X16468. 1		12	4. 8	376	2. 57	0. 62	0. 41
463	Trachea	205941_s_at	collagen, type X, alpha 1 (Schmid metaphyseal c	Hs. 179729. 0	gb:AI376003_RC		6	3. 6	173	2. 24	1. 38	0. 24
464	Trachea	206905_s_at	matrilin 1, cartilage matrix protein	Hs. 150366. 0	gb:NM_002379. 2	MATN1	1	3. 6	318	2. 50	1. 12	0. 38
465	Trachea	207913_at	cytochrome P450, subfamily IIF, polypeptide 1	Hs. 72913. 0	gb:NM_000774. 1	CYP2F1	19	3. 6	84	1. 93	0. 74	0. 33
466	Trachea	209720_s_at	serine (or cysteine) proteinase inhibitor, cla	Hs. 227948. 0	gb:BC005224. 1		18	3. 3	456	2. 66	0. 78	0. 57
467	Trachea	205749_at	cytochrome P450, subfamily I (aromatic compound	Hs. 72912. 0	gb:NM_000499. 2	CYP1A1	15	3. 1	258	2. 41	1. 58	0. 27
468	Lung	205982_x_at	surfactant, pulmonary-associated protein C	Hs. 1074. 0	gb:NM_003018. 1	SFTPC		24. 9	6809	3. 83	0. 82	0. 12
469	Lung	214387_x_at	surfactant, pulmonary-associated protein C	Hs. 1074. 1	gb:AA633841_RC			16	6358	3. 80	0. 98	0. 18
470	Lung	211735_x_at			gb:BC005913. 1			15. 5	6770	3. 83	0. 84	0. 19
471	Lung	214199_at	surfactant, pulmonary-associated protein D	Hs. 253495. 0	gb:NM_003019. 1	SFTPD	10	12. 7	1072	3. 03	0. 54	0. 20
472	Lung	215454_x_at	surfactant, pulmonary-associated protein C	Hs. 1074. 3	gb:AI831055_RC			12. 1	1918	3. 28	0. 49	0. 23
473	Lung	213936_x_at	N-acylsphingosine amidohydrolase (acid ceramida	Hs. 264330. 1	gb:AW276646_RC		2	10. 4	2201	3. 34	1. 82	0. 15
474	Lung	37004_at						7. 6	3736	3. 57	1. 00	0. 34
475	Lung	38691_s_at						7. 5	6282	3. 80	1. 84	0. 26
476	Lung	207442_at	colony stimulating factor 3 (granulocyte)	Hs. 2233. 0	gb:NM_000759. 1	CSF3	17	7. 3	111	2. 05	0. 76	0. 18
477	Lung	209810_at	surfactant, pulmonary-associated protein B	Hs. 76305. 0	gb:J02761. 1	SFTP3	2	6. 8	3975	3. 60	1. 16	0. 36
478	Lung	217046_s_at	advanced glycosylation end product-specific rec	Hs. 184. 1	gb:AJ133822. 1	RAGEsec	6	6. 8	346	2. 54	0. 48	0. 30
479	Lung	204811_s_at	calcium channel, voltage-dependent, alpha 2del	Hs. 127436. 0	gb:NM_006030. 1	CACNA2D2	3	6. 1	429	2. 63	1. 69	0. 15
480	Lung	214354_x_at	N-acylsphingosine amidohydrolase (acid ceramida	Hs. 264330. 0	gb:T91506_RC		2	6	2407	3. 38	1. 85	0. 25
481	Lung	209959_at	nuclear receptor subfamily 4, group A, member 3	Hs. 80561. 1	gb:U12767. 1	MINOR	9	5. 2	258	2. 41	1. 26	0. 22
482	Lung	210081_at	advanced glycosylation end product-specific rec	Hs. 184. 0	gb:AB036432. 1	RAGE	6	5	805	2. 91	1. 24	0. 34
483	Lung	214974_x_at	Homo sapiens cDNA: FLJ22893 fis, clone KAT04792	Hs. 287716. 0	gb:AK026546. 1		4	4. 6	194	2. 29	1. 16	0. 25
484	Lung	205569_at	similar to lysosome-associated membrane glycopr	Hs. 10887. 0	gb:NM_014398. 1	TSC403	3	4. 5	1665	3. 22	1. 45	0. 39
485	Lung	207316_at	hyaluronan synthase 1	Hs. 57697. 0	gb:NM_001523. 1	HAS1	19	4. 1	247	2. 39	1. 14	0. 30
486	Lung	219795_at	solute carrier family 6 (neurotransmitter trans	Hs. 162211. 0	gb:NM_007231. 1	SLC6A14	X	3. 8	381	2. 58	1. 22	0. 36
487	Lung	220677_s_at	a disintegrin-like and metalloprotease (reprol	Hs. 271605. 0	gb:NM_007037. 1	ADAMTS8	11	3. 6	143	2. 16	0. 97	0. 33
488	Lung	215059_at	Homo sapiens mRNA; cDNA DKFZp564G112 (from clor	Hs. 51515. 0	gb:AA053967_RC		4	3. 4	131	2. 12	0. 63	0. 44
489	Lung	206026_s_at	tumor necrosis factor, alpha-induced protein 6	Hs. 29352. 0	gb:NM_007115. 1	TNFAIP6	2	3. 2	239	2. 38	1. 64	0. 23
490	Lung	219725_at	triggering receptor expressed on myeloid cells	Hs. 44234. 0	gb:NM_018965. 1	TREM2	6	2. 9	238	2. 38	1. 31	0. 37
491	Lung	218995_s_at	endothelin 1	Hs. 2271. 0	gb:NM_001955. 1	EDN1	6	2. 8	420	2. 62	1. 55	0. 38
492	Breast	208241_at	neuregulin 1	Hs. 172816. 4	gb:NM_004495. 1	NRG1		5. 5	195	2. 29	1. 78	0. 09
493	Breast	215271_at	hypothetical protein similar to tenascin-R	Hs. 156369. 0	gb:BF432086_RC		1	5. 3	181	2. 26	1. 45	0. 15
494	Breast	206825_at	oxytocin receptor	Hs. 2820. 0	gb:NM_000916. 2	OXTR	3	4. 9	479	2. 68	1. 52	0. 24

Fig. 54

	A	B	C	D	E	F	G	H	I	J	K	L
495	Breast	212960_at	KIAA0882 protein	Hs. 90419. 0	gb:BE646554_RC		4	4. 8	267	2. 43	1. 80	0. 13
496	Breast	204694_at	alpha-fetoprotein	Hs. 155421. 0	gb:NM_001134. 1	AFP	4	4. 4	174	2. 24	1. 30	0. 21
497	Breast	202035_s_at	secreted frizzled-related protein 1	Hs. 7306. 0	gb:AI332407_RC		8	4. 2	177	2. 25	1. 18	0. 25
498	Breast	204851_s_at	doublecortex; lissencephaly, X-linked (doublec	Hs. 34780. 0	gb:AF040254. 1	DCX	X	3. 9	100	2. 00	0. 56	0. 37
499	Breast	209884_s_at	solute carrier family 4, sodium bicarbonate co	Hs. 132904. 1	gb:AF047033. 1	SLC4A7	3	3. 9	241	2. 38	1. 54	0. 22
500	Breast	218502_s_at	trichorhinophalangeal syndrome I	Hs. 26102. 0	gb:NM_014112. 1	TRPS1	8	3. 9	314	2. 50	1. 60	0. 23
501	Breast	219768_at	hypothetical protein FLJ22418	Hs. 36563. 0	gb:NM_024626. 1	FLJ22418	1	3. 9	399	2. 60	1. 35	0. 32
502	Breast	203438_at	stanniocalcin 2	Hs. 155223. 0	gb:AI435828_RC		5	3. 5	454	2. 66	1. 36	0. 37
503	Breast	211277_x_at	amyloid beta (A4) precursor protein (protease r	Hs. 177486. 1	gb:BC004369. 1		21	3. 4	382	2. 58	1. 84	0. 22
504	Breast	211466_at	nuclear factor IB	Hs. 33287. 0	gb:U70862. 1		9	3. 2	126	2. 10	0. 71	0. 43
505	Breast	211756_at			gb:BC005961. 1		12	1. 8	58	1. 76	0. 88	0. 48
506	Skin	206969_at	keratin, hair, acidic, 4	Hs. 296942. 0	gb:NM_021013. 2	KRTHA4	17	14. 8	1318	3. 12	0. 76	0. 16
507	Skin	208483_x_at	keratin, hair, acidic, 3A	Hs. 197874. 0	gb:NM_004138. 1	KRTHA3A	17	14. 5	2182	3. 34	0. 42	0. 20
508	Skin	207787_at	keratin, hair, acidic, 3B	Hs. 32950. 0	gb:NM_002279. 2	KRTHA3B	17	13. 3	917	2. 96	0. 91	0. 16
509	Skin	220970_s_at			gb:NM_030977. 1	KRTAP2. 1A	17	12. 1	1369	3. 14	0. 82	0. 19
510	Skin	206677_at	keratin, hair, acidic, 1	Hs. 41696. 0	gb:NM_002277. 1	KRTHA1	17	11. 2	2428	3. 39	1. 45	0. 17
511	Skin	215189_at	keratin, hair, basic, 6 (monilethrix)	Hs. 278658. 0	gb:X99142. 1	hHb6	12	11. 1	2028	3. 31	1. 83	0. 13
512	Skin	206192_at	corneodesmosin	Hs. 507. 0	gb:L20815. 1		6	10. 7	323	2. 51	0. 95	0. 14
513	Skin	220635_at	SPR1 protein	Hs. 146824. 0	gb:NM_014069. 1	SPR1	6	10. 1	616	2. 79	1. 10	0. 17
514	Skin	206193_s_at	corneodesmosin	Hs. 507. 0	gb:NM_001264. 1	CDSN	6	9. 9	362	2. 56	0. 86	0. 17
515	Skin	206033_s_at	desmocollin 3	Hs. 41690. 0	gb:NM_001941. 2	DSC3	18	9. 4	897	2. 95	1. 55	0. 15
516	Skin	221854_at	ESTs	Hs. 292911. 0	gb:AI378979_RC		1	9	1001	3. 00	0. 64	0. 26
517	Skin	207669_at	keratin, hair, basic, 3	Hs. 182506. 0	gb:NM_002282. 1	KRTHB3	12	8. 5	884	2. 95	0. 83	0. 25
518	Skin	206400_at	lectin, galactoside-binding, soluble, 7 (galect	Hs. 99923. 0	gb:NM_002307. 1	LGALS7	19	7. 9	3294	3. 52	0. 81	0. 34
519	Skin	220972_s_at			gb:NM_030975. 1	KRTAP9. 9	17	7. 2	855	2. 93	0. 30	0. 37
520	Skin	216921_s_at	keratin, hair, acidic, 5	Hs. 73082. 1	gb:X90763. 1	hHa5	17	7. 1	1683	3. 23	1. 40	0. 26
521	Skin	216810_at	Homo sapiens mRNA for keratin associated protei	Hs. 307020. 0	gb:AJ406939. 1	KRTAP4. 7	17	7	1262	3. 10	1. 18	0. 27
522	Skin	207670_at	keratin, hair, basic, 5	Hs. 182507. 0	gb:NM_002283. 1	KRTHB5	12	6. 8	2262	3. 35	1. 70	0. 24
523	Skin	207720_at	loricrin	Hs. 251680. 0	gb:NM_000427. 1	LOR	1	6. 8	3507	3. 54	0. 89	0. 39
524	Skin	206032_at	desmocollin 3	Hs. 41690. 0	gb:AI797281_RC		18	6. 6	1053	3. 02	1. 64	0. 21
525	Skin	213780_at	trichohyalin	Hs. 82276. 0	gb:N30878_RC		1	6. 6	1377	3. 14	1. 41	0. 26
526	Skin	206156_at	gap junction protein, beta 5 (connexin 31. 1)	Hs. 198249. 0	gb:NM_005268. 1	GJB5	1	6. 4	93	1. 97	0. 89	0. 17
527	Skin	206165_s_at	chloride channel, calcium activated, family mer	Hs. 241551. 0	gb:NM_006536. 2	CLCA2	1	6. 4	742	2. 87	1. 60	0. 20
528	Skin	220978_at			gb:NM_030966. 1	KRTAP1. 3	17	6. 3	1474	3. 17	0. 80	0. 38
529	Skin	221297_at	G protein-coupled receptor, family C, group 5,	Hs. 283073. 0	gb:NM_018654. 1	GPRC5D	12	6. 3	558	2. 75	2. 02	0. 11
530	Skin	205338_s_at	dopachrome tautomerase (dopachrome delta-isomer	Hs. 301865. 0	gb:NM_001922. 2	DCT	13	6. 1	360	2. 56	1. 90	0. 11
531	Skin	206427_s_at	melan-A	Hs. 154069. 0	gb:U06654. 1		9	6. 1	208	2. 32	1. 58	0. 12
532	Skin	220013_at	hypothetical protein FLJ22408	Hs. 156457. 0	gb:NM_024794. 1	FLJ22408	19	6	253	2. 40	0. 74	0. 28

Fig. 55

	A	B	C	D	E	F	G	H	I	J	K	L
533	Skin	213711_at	keratin, hair, basic, 1	Hs. 32952. 0	gb:NM_002281. 1	KRTHB1	12	5. 9	625	2. 80	0. 66	0. 36
534	Skin	207324_s_at	desmocollin 1	Hs. 69752. 0	gb:NM_004948. 2	DSC1	18	5. 8	1523	3. 18	1. 17	0. 35
535	Skin	220976_s_at			gb:NM_030967. 1	KRTAP1. 1	17	5. 8	1339	3. 13	1. 47	0. 29
536	Skin	209800_at	keratin 16 (focal non-epidermolytic palmoplanta	Hs. 115947. 0	gb:AF061812. 1	KRT16A	17	5. 6	960	2. 98	0. 58	0. 43
537	Skin	217395_at	Human metallothionein IV (MTIV) gene, complete	Hs. 194762. 0	gb:X55503			5. 5	239	2. 38	1. 01	0. 25
538	Skin	210576_at	cytochrome P450, subfamily IVF, polypeptide 8	Hs. 268554. 0	gb:AF133298. 1	CYP4F8	19	5. 3	123	2. 09	0. 25	0. 35
539	Skin	206276_at	lymphocyte antigen 6 complex, locus D	Hs. 3185. 0	gb:NM_003695. 1	E48	8	5. 2	719	2. 86	0. 67	0. 42
540	Skin	215704_at	filaggrin	Hs. 73995. 0	gb:AL356504		1	5. 1	1564	3. 19	0. 78	0. 47
541	Skin	207114_at	G6C protein	Hs. 241586. 0	gb:NM_025261. 1	G6C	6	4. 9	1017	3. 01	1. 69	0. 27
542	Skin	206166_s_at	chloride channel, calcium activated, family mer	Hs. 241551. 0	gb:AF043977. 1	hCLCA2	1	4. 8	294	2. 47	1. 20	0. 26
543	Skin	207710_at	skin-specific protein	Hs. 234766. 0	gb:NM_014357. 1	XP5	1	4. 8	502	2. 70	1. 12	0. 33
544	Skin	209260_at	stratifin	Hs. 184510. 0	gb:BC000329. 1		1	4. 8	948	2. 98	0. 97	0. 42
545	Skin	205337_at	dopachrome tautomerase (dopachrome delta-isomer	Hs. 301865. 0	gb:AL139318		13	4. 7	427	2. 63	1. 07	0. 33
546	Skin	216512_s_at	dopachrome tautomerase (dopachrome delta-isomer	Hs. 301865. 1	gb:AL139318		13	4. 7	184	2. 26	1. 37	0. 19
547	Skin	221470_s_at	Interleukin-1 Superfamily z	Hs. 166371. 1	gb:NM_014439. 1	FIL1(ZETA)	2	4. 7	67	1. 82	0. 42	0. 30
548	Skin	205778_at	kallikrein 7 (chymotryptic, stratum corneum)	Hs. 151254. 0	gb:NM_005046. 1	KLK7	19	4. 6	426	2. 63	1. 57	0. 23
549	Skin	205490_x_at	gap junction protein, beta 3, 31kD (connexin 31	Hs. 98485. 0	gb:BF060667_RC		1	4. 5	435	2. 64	1. 45	0. 27
550	Skin	205916_at	S100 calcium-binding protein A7 (psoriasin 1)	Hs. 112408. 0	gb:NM_002963. 2	S100A7	10	4. 4	659	2. 82	0. 77	0. 46
551	Skin	209848_s_at	silver (mouse homolog) like	Hs. 95972. 0	gb:U01874. 1		12	4. 4	251	2. 40	1. 69	0. 16
552	Skin	206642_at	desmoglein 1	Hs. 2633. 0	gb:NM_001942. 1	DSG1	18	4. 3	904	2. 96	0. 86	0. 49
553	Skin	207457_s_at	megakaryocyte-enhanced gene transcript 1 protei	Hs. 241587. 0	gb:NM_021246. 1	MEGT1	6	4. 1	339	2. 53	1. 44	0. 27
554	Skin	219832_s_at	homeo box C13	Hs. 118608. 0	gb:NM_017410. 1	HOXC13	12	4. 1	104	2. 02	0. 84	0. 29
555	Skin	206164_at	chloride channel, calcium activated, family mer	Hs. 241551. 0	gb:NM_006536. 2	CLCA2	1	3. 9	209	2. 32	0. 68	0. 42
556	Skin	206884_s_at	sciellin	Hs. 115166. 0	gb:NM_003843. 1	SCEL	13	3. 9	489	2. 69	0. 60	0. 53
557	Skin	206027_at	S100 calcium-binding protein A3	Hs. 2961. 0	gb:NM_002960. 1	S100A3	1	3. 7	404	2. 61	1. 30	0. 36
558	Skin	214536_at	ARS component B	Hs. 103505. 0	gb:NM_020427. 1	ARS	8	3. 7	252	2. 40	1. 18	0. 33
559	Skin	217528_at	ESTs	Hs. 72365. 0	gb:BF003134_RC			3. 7	333	2. 52	0. 77	0. 47
560	Skin	213154_s_at	KIAA0699 protein	Hs. 17411. 1	gb:AI934125_RC		9	3. 6	432	2. 63	1. 94	0. 19
561	Skin	222223_s_at	interleukin-1 receptor antagonist homolog 1	Hs. 207224. 1	gb:AF216693			3. 6	137	2. 14	1. 48	0. 18
562	Skin	208539_x_at	small proline-rich protein 2B	Hs. 231622. 0	gb:NM_006945. 1	SPRR2B	1	3. 5	233	2. 37	1. 04	0. 38
563	Skin	214240_at	galanin	Hs. 1907. 0	gb:AL556409		11	3. 5	199	2. 30	1. 49	0. 23
564	Skin	220403_s_at	p53-regulated apoptosis-inducing protein 1	Hs. 160953. 0	gb:NM_022112. 1	P53AIP1	11	3. 5	46	1. 66	0. 50	0. 33
565	Skin	202179_at	bleomycin hydrolase	Hs. 78943. 0	gb:NM_000386. 1	BLMH	17	3. 4	328	2. 52	1. 77	0. 22
566	Skin	205822_s_at	3-hydroxy-3-methylglutaryl-Coenzyme A synthase	Hs. 77910. 0	gb:NM_002130. 1	HMGCS1	5	3. 4	376	2. 58	1. 71	0. 25
567	Skin	206023_at	neuromedin U	Hs. 2841. 0	gb:NM_006681. 1	NMU	4	3. 4	142	2. 15	0. 62	0. 45
568	Skin	207065_at	cytokeratin type II	Hs. 145949. 0	gb:NM_004693. 1	K6HF	12	3. 4	311	2. 49	1. 51	0. 29
569	Skin	207146_at	keratin, hair, acidic, 2	Hs. 41752. 0	gb:NM_002278. 1	KRTHA2	17	3. 4	248	2. 39	1. 65	0. 22
570	Skin	220620_at	NICE-1 protein	Hs. 110196. 0	gb:NM_019060. 1	NICE-1	1	3. 4	214	2. 33	1. 23	0. 32

Fig. 56

	A	B	C	D	E	F	G	H	I	J	K	L
571	Skin	206605_at	protease, serine, 22	Hs. 997. 0	gb:NM_006025. 1	P11	12	3. 3	179	2. 25	1. 24	0. 30
572	Skin	207955_at	small inducible cytokine subfamily A (Cys-Cys),	Hs. 225948. 0	gb:NM_006664. 1	SCYA27	9	3. 3	123	2. 09	1. 09	0. 30
573	Skin	215465_at	DKFZP434G232 protein	Hs. 134585. 0	gb:AL080207. 1	DKFZp434G2	2	3. 3	148	2. 17	1. 07	0. 33
574	Skin	209098_s_at	jagged 1 (Alagille syndrome)	Hs. 91143. 0	gb:U61276. 1	HJ1	20	3. 2	267	2. 43	1. 71	0. 22
575	Skin	206426_at	melan-A	Hs. 154069. 0	gb:NM_005511. 1	MLANA	9	3. 1	86	1. 93	1. 03	0. 30
576	Skin	207109_at	POU transcription factor	Hs. 227115. 0	gb:NM_014352. 1	11. Oct	11	3. 1	85	1. 93	0. 61	0. 42
577	Skin	216935_at	Homo sapiens skin-specific protein (xp33) mRNA,	Hs. 113261. 0	gb:AF005082. 1	xp33	1	2. 7	83	1. 92	1. 00	0. 34
578	Skeltal musc	207284_s_at	aspartate beta-hydroxylase	Hs. 283664. 1	gb:NM_020164. 1	LOC56921	8	8. 3	138	2. 14	0. 39	0. 21
579	Skeltal musc	206633_at	cholinergic receptor, nicotinic, alpha polypept	Hs. 2266. 0	gb:NM_000079. 1	CHRNA1	2	7. 7	821	2. 91	1. 58	0. 17
580	Skeltal musc	206121_at	adenosine monophosphate deaminase 1 (isoform M)	Hs. 89570. 0	gb:NM_000036. 1	AMPD1	1	7. 6	2369	3. 37	1. 65	0. 23
581	Skeltal musc	206981_at	sodium channel, voltage-gated, type IV, alpha p	Hs. 46038. 0	gb:NM_000334. 1	SCN4A	17	6. 6	278	2. 44	0. 62	0. 28
582	Skeltal musc	202417_at	Kelch-like ECH-associated protein 1	Hs. 57729. 0	gb:NM_012289. 1	KIAA0132	19	6. 2	543	2. 73	2. 00	0. 12
583	Skeltal musc	206844_at	fructose-1, 6-bisphosphatase 2	Hs. 61255. 0	gb:NM_003837. 1	FBP2	9	6. 2	165	2. 22	0. 64	0. 25
584	Skeltal musc	208534_s_at	GTPase activating protein-like	Hs. 184367. 1	gb:NM_006989. 1	GAPL	7	6. 2	243	2. 38	0. 82	0. 25
585	Skeltal musc	213567_at	Homo sapiens clone 23728 mRNA sequence	Hs. 153106. 0	gb:BF431965_RC		3	6. 2	604	2. 78	1. 91	0. 14
586	Skeltal musc	210632_s_at	sarcoglycan, alpha (50kD dystrophin-associated	Hs. 99931. 1	gb:L35853. 1		17	5. 9	256	2. 41	0. 49	0. 33
587	Skeltal musc	218177_at	CHMP1. 5 protein	Hs. 42733. 0	gb:AA293502		18	5. 9	566	2. 75	1. 87	0. 15
588	Skeltal musc	201388_at	proteasome (prosome, macropain) 26S subunit, n	Hs. 9736. 0	gb:NM_002809. 1	PSMD3	17	5. 8	464	2. 67	2. 02	0. 11
589	Skeltal musc	205951_at	myosin, heavy polypeptide 1, skeletal muscle, a	Hs. 231581. 0	gb:NM_005963. 2	MYH1	17	5. 8	3279	3. 52	0. 76	0. 47
590	Skeltal musc	206891_at	actinin, alpha 3	Hs. 1216. 0	gb:NM_001104. 1	ACTN3	11	5. 8	864	2. 94	1. 55	0. 24
591	Skeltal musc	205693_at	troponin T3, skeletal, fast	Hs. 73454. 0	gb:NM_006757. 1	TNNT3	11	5. 7	1393	3. 14	1. 51	0. 29
592	Skeltal musc	37950_at						5. 4	129	2. 11	1. 56	0. 10
593	Skeltal musc	210837_s_at	phosphodiesterase 4D, cAMP-specific (dunce (Dr	Hs. 172081. 3	gb:AF012074. 1	PDE4D	5	5. 2	111	2. 05	1. 24	0. 16
594	Skeltal musc	206996_x_at	calcium channel, voltage-dependent, beta 1 subu	Hs. 635. 0	gb:NM_000723. 1	CACNB1	17	5. 1	598	2. 78	1. 55	0. 24
595	Skeltal musc	205330_at	meningioma (disrupted in balanced translocation	Hs. 268515. 0	gb:NM_002430. 1	MN1	22	5	419	2. 62	1. 83	0. 16
596	Skeltal musc	206895_at	protein phosphatase 1, regulatory (inhibitor) s	Hs. 127614. 0	gb:NM_002711. 1	PPP1R3		5	606	2. 78	0. 80	0. 40
597	Skeltal musc	218841_at	hypothetical protein MGC5540	Hs. 267400. 0	gb:NM_024095. 1	MGC5540		5	374	2. 57	1. 52	0. 21
598	Skeltal musc	208264_s_at	eukaryotic translation initiation factor 3, sub	Hs. 173987. 0	gb:NM_003758. 1	EIF3S1	15	4. 9	280	2. 45	1. 77	0. 14
599	Skeltal musc	219827_at	uncoupling protein 3 (mitochondrial, proton car	Hs. 101337. 0	gb:NM_003356. 2	UCP3	11	4. 9	358	2. 55	0. 94	0. 33
600	Skeltal musc	208148_at	myosin, heavy polypeptide 4, skeletal muscle	Hs. 272207. 0	gb:NM_017533. 1	MYH4	17	4. 8	165	2. 22	0. 45	0. 37
601	Skeltal musc	209533_s_at	phospholipase A2-activating protein	Hs. 27182. 0	gb:AF145020. 1	PLA2P	9	4. 7	194	2. 29	1. 77	0. 11
602	Skeltal musc	210836_x_at	phosphodiesterase 4D, cAMP-specific (dunce (Dr	Hs. 172081. 2	gb:AF012073. 1	PDE4D	5	4. 7	145	2. 16	1. 51	0. 14
603	Skeltal musc	213349_at	KIAA0779 protein	Hs. 179507. 0	gb:AI934469_RC		3	4. 7	315	2. 50	1. 80	0. 15
604	Skeltal musc	221276_s_at			gb:NM_030786. 1	SYNCOILIN	1	4. 7	437	2. 64	1. 48	0. 25
605	Skeltal musc	214099_s_at	Homo sapiens cDNA FLJ10757 fis, clone NT2RP300	Hs. 323580. 0	gb:AK001619. 1		1	4. 5	215	2. 33	1. 04	0. 29
606	Skeltal musc	205231_s_at	epilepsy, progressive myoclonus type 2, Lafora	Hs. 22464. 0	gb:NM_005670. 1	EPM2A	6	4. 4	222	2. 35	1. 48	0. 20
607	Skeltal musc	207349_s_at	uncoupling protein 3 (mitochondrial, proton car	Hs. 101337. 1	gb:NM_022803. 1	UCP3	11	4. 4	509	2. 71	1. 12	0. 36
608	Skeltal musc	208567_s_at	potassium inwardly-rectifying channel, subfamil	Hs. 248143. 0	gb:NM_002244. 1	KCNJN1	17	4. 4	101	2. 00	0. 66	0. 30

Fig. 57

	A	B	C	D	E	F	G	H	I	J	K	L
609	Skeltal musc	213741_s_at	karyopherin alpha 1 (importin alpha 5)	Hs. 169149. 1	gb:BF575685		3	4. 4	203	2. 31	1. 77	0. 12
610	Skeltal musc	217097_s_at	hypothetical protein DKFZp564F013	Hs. 128653. 1	gb:AC004990			4. 4	248	2. 39	1. 41	0. 22
611	Skeltal musc	222287_at	ESTs	Hs. 291232. 0	gb:AW969675		6	4. 4	887	2. 95	0. 38	0. 58
612	Skeltal musc	212707_s_at	GTPase activating protein-like	Hs. 184367. 0	gb:AI738591_RC		7	4. 2	208	2. 32	1. 41	0. 22
613	Skeltal musc	210967_x_at	calcium channel, voltage-dependent, beta 1 subu	Hs. 635. 1	gb:M92301. 1		17	4	561	2. 75	1. 62	0. 28
614	Skeltal musc	205120_s_at	sarcoglycan, beta (43kD dystrophin-associated g	Hs. 77501. 0	gb:U29586. 1		4	3. 9	382	2. 58	1. 83	0. 19
615	Skeltal musc	206106_at	mitogen-activated protein kinase 12	Hs. 55039. 0	gb:AL022328		22	3. 9	344	2. 54	1. 24	0. 33
616	Skeltal musc	206612_at	calcium channel, voltage-dependent, gamma subur	Hs. 147989. 0	gb:NM_000727. 1	CACNG1	17	3. 9	366	2. 56	1. 03	0. 40
617	Skeltal musc	217924_at	hypothetical protein FLJ22195	Hs. 25999. 0	gb:AL523965		6	3. 9	226	2. 35	1. 55	0. 21
618	Skeltal musc	205377_s_at	acetylcholinesterase (YT blood group)	Hs. 154495. 0	gb:AI190022_RC		7	3. 8	181	2. 26	1. 24	0. 27
619	Skeltal musc	206703_at	cholinergic receptor, nicotinic, beta polypepti	Hs. 89739. 0	gb:NM_000747. 1	CHRNB1	NA	3. 8	285	2. 45	1. 63	0. 22
620	Skeltal musc	211764_s_at			gb:BC005980. 1		10	3. 8	326	2. 51	1. 85	0. 17
621	Skeltal musc	214590_s_at	ubiquitin-conjugating enzyme E2D 1 (homologous	Hs. 318435. 0	gb:AL545760		10	3. 8	88	1. 95	1. 06	0. 24
622	Skeltal musc	207145_at	growth differentiation factor 8	Hs. 41565. 0	gb:NM_005259. 1	GDF8	2	3. 7	208	2. 32	0. 12	0. 60
623	Skeltal musc	207968_s_at	MADS box transcription enhancer factor 2, polyp	Hs. 78995. 0	gb:NM_002397. 1	MEF2C		3. 7	265	2. 42	1. 53	0. 24
624	Skeltal musc	211890_x_at	calpain 3, (p94)	Hs. 40300. 4	gb:AF127765. 3	CAPN3	15	3. 7	844	2. 93	0. 94	0. 54
625	Skeltal musc	201878_at	ariadne (Drosophila) homolog, ubiquitin-conjuga	Hs. 181461. 0	gb:N25546_RC		15	3. 6	257	2. 41	1. 68	0. 20
626	Skeltal musc	214145_s_at	KIAA0599 protein	Hs. 198037. 1	gb:BG223341_RC		14	3. 6	326	2. 51	0. 93	0. 44
627	Skeltal musc	202056_at	karyopherin alpha 1 (importin alpha 5)	Hs. 169149. 0	gb:AW051311_RC		3	3. 5	352	2. 55	1. 78	0. 22
628	Skeltal musc	204498_s_at	adenylate cyclase 9	Hs. 20196. 0	gb:NM_001116. 1	ADCY9	16	3. 5	353	2. 55	1. 84	0. 21
629	Skeltal musc	204647_at	Homer, neuronal immediate early gene, 3	Hs. 166146. 0	gb:NM_004838. 1	HOMER-3	19	3. 5	310	2. 49	1. 69	0. 23
630	Skeltal musc	209244_s_at	KIAA0706 gene product	Hs. 139648. 0	gb:BE885926		17	3. 5	230	2. 36	1. 21	0. 33
631	Skeltal musc	216088_s_at	proteasome (prosome, macropain) subunit, alpha	Hs. 233952. 3	gb:AL078633		20	3. 5	243	2. 39	1. 85	0. 15
632	Skeltal musc	204217_s_at	reticulon 2	Hs. 3803. 0	gb:NM_005619. 1	RTN2	19	3. 4	399	2. 60	1. 26	0. 40
633	Skeltal musc	211273_s_at	T-box 1	Hs. 173984. 1	gb:AF012130. 1	TBX1	22	3. 4	223	2. 35	0. 62	0. 51
634	Skeltal musc	214846_s_at	KIAA1330 protein	Hs. 301242. 0	gb:AB037751. 1	KIAA1330	15	3. 4	208	2. 32	1. 30	0. 30
635	Skeltal musc	217515_s_at	Homo sapiens calcium channel, voltage-dependent	Hs. 328199. 0	gb:AI391509_RC		1	3. 3	294	2. 47	1. 24	0. 37
636	Skeltal musc	205378_s_at	acetylcholinesterase (YT blood group)	Hs. 154495. 0	gb:NM_015831. 1	ACHE	7	3. 2	151	2. 18	0. 79	0. 43
637	Skeltal musc	207110_at	potassium inwardly-rectifying channel, subfami	Hs. 2363. 0	gb:NM_021012. 1	KCNJ12	17	3. 2	136	2. 13	0. 86	0. 40
638	Skeltal musc	207558_s_at	paired-like homeodomain transcription factor 2	Hs. 92282. 0	gb:NM_000325. 1	PITX2	4	3. 2	172	2. 24	0. 88	0. 43
639	Skeltal musc	210870_s_at	epilepsy, progressive myoclonus type 2, Lafora	Hs. 22464. 1	gb:BC005286. 1		6	3. 2	164	2. 22	1. 53	0. 22
640	Skeltal musc	212427_at	KIAA0368 protein	Hs. 3852. 1	gb:AB002366. 1	KIAA0368	9	3. 2	158	2. 20	1. 35	0. 26
641	Skeltal musc	213955_at	Homo sapiens clone B18 unknown mRNA	Hs. 91626. 0	gb:BE673151_RC		5	3. 2	316	2. 50	1. 48	0. 32
642	Skeltal musc	215286_s_at	hypothetical protein DKFZp564F013	Hs. 128653. 2	gb:AL050389. 1		7	3. 2	84	1. 93	1. 23	0. 22
643	Skeltal musc	202968_s_at	dual-specificity tyrosine-(Y)-phosphorylation r	Hs. 173135. 1	gb:Y09216. 1	Dyrk2	12	3. 1	260	2. 41	1. 47	0. 30
644	Skeltal musc	203138_at	histone acetyltransferase 1	Hs. 13340. 0	gb:NM_003642. 1	HAT1	2	3. 1	294	2. 47	1. 87	0. 20
645	Skeltal musc	214121_x_at	enigma (LIM domain protein)	Hs. 102948. 1	gb:AA086229_RC		5	3. 1	540	2. 73	1. 01	0. 55
646	Skeltal musc	218744_s_at	protein kinase C and casein kinase substrate in	Hs. 256310. 0	gb:NM_016223. 1	LOC51165	11	3. 1	208	2. 32	0. 85	0. 47

Fig. 58

	A	B	C	D	E	F	G	H	I	J	K	L
647	Skeltal musc	204058_at	malic enzyme 1, NADP(+)-dependent, cytosolic	Hs. 14732. 0	gb:AL049699		6	3	388	2. 59	1. 65	0. 31
648	Skeltal musc	214475_x_at	calpain 3, (p94)	Hs. 40300. 3	gb:AF127764. 1	CAPN3	15	3	580	2. 76	0. 82	0. 64
649	Skeltal musc	208208_at	myosin, heavy polypeptide 13, skeletal muscle	Hs. 278488. 0	gb:NM_003802. 1	MYH13	17	2. 4	114	2. 06	1. 13	0. 39
650	Skeltal musc	212797_at	sortilin 1	Hs. 281706. 1	gb:BE742268		1	2	93	1. 97	1. 26	0. 36
651	Heart	219942_at	myosin light chain 2a	Hs. 75636. 0	gb:NM_021223. 1	LOC58498	7	14	3033	3. 48	0. 81	0. 19
652	Heart	205742_at	troponin I, cardiac	Hs. 54668. 0	gb:NM_000363. 1	TNNI3	18	10. 4	5404	3. 73	1. 47	0. 22
653	Heart	206801_at	natriuretic peptide precursor B	Hs. 219140. 0	gb:NM_002521. 1	NPPB	1	8. 3	1446	3. 16	1. 32	0. 22
654	Heart	208040_s_at			gb:NM_000256. 2	MYBPC3	11	7. 7	2126	3. 33	1. 52	0. 23
655	Heart	215389_s_at	troponin T2, cardiac	Hs. 296865. 1	gb:X79857. 1		1	7. 4	3028	3. 48	1. 23	0. 30
656	Heart	217359_s_at	neural cell adhesion molecule 1	Hs. 167988. 3	gb:M22094. 1	NCAM	11	7. 4	472	2. 67	1. 80	0. 12
657	Heart	205741_s_at	dystrobrevin, alpha	Hs. 54435. 0	gb:NM_001392. 1	DTNA	18	6. 3	415	2. 62	1. 64	0. 16
658	Heart	219647_at	popeye protein 2	Hs. 77208. 0	gb:NM_022135. 1	POP2	3	5. 9	493	2. 69	1. 58	0. 19
659	Heart	219829_at	integrin beta 1 binding protein (melusin) 2	Hs. 109999. 0	gb:NM_012278. 1	ITGB1BP2	X	5. 6	512	2. 71	1. 03	0. 30
660	Heart	207303_at	phosphodiesterase 1C, calmodulin-dependent (70	Hs. 41718. 0	gb:NM_005020. 1	PDE1C	7	5. 2	249	2. 40	1. 71	0. 13
661	Heart	205918_at	solute carrier family 4, anion exchanger, membe	Hs. 1176. 0	gb:NM_005070. 1	SLC4A3	2	4. 8	540	2. 73	0. 67	0. 42
662	Heart	201025_at	KIAA0741 gene product	Hs. 158688. 0	gb:AB018284. 1	KIAA0741	2	4. 7	238	2. 38	1. 86	0. 11
663	Heart	214044_at	Homo sapiens ryanodine receptor 2 (cardiac) (R	Hs. 332699. 0	gb:BE968750		1	4. 6	742	2. 87	1. 20	0. 37
664	Heart	207557_s_at	ryanodine receptor 2 (cardiac)	Hs. 90821. 0	gb:NM_001035. 1	RYR2	1	4. 6	523	2. 72	0. 70	0. 44
665	Heart	206578_at	cardiac-specific homeo box	Hs. 54473. 0	gb:NM_004387. 1	CSX	5	4. 6	370	2. 57	1. 15	0. 31
666	Heart	219338_s_at	hypothetical protein FLJ20156	Hs. 12692. 0	gb:NM_017691. 1	FLJ20156	15	4. 6	227	2. 36	1. 71	0. 14
667	Heart	211476_at	calcineurin-binding protein calsarcin-1	Hs. 42346. 1	gb:AY013295. 1			4. 3	725	2. 86	0. 48	0. 56
668	Heart	209957_s_at	natriuretic peptide precursor A	Hs. 75640. 0	gb:M30262. 1		1	4. 2	416	2. 62	1. 40	0. 29
669	Heart	207413_s_at	sodium channel, voltage-gated, type V, alpha p	Hs. 169331. 0	gb:NM_000335. 1	SCN5A		4. 1	210	2. 32	1. 34	0. 24
670	Heart	215918_s_at	spectrin, beta, non-erythrocytic 1	Hs. 107164. 4	gb:AA131826		2	4	434	2. 64	1. 75	0. 23
671	Heart	214285_at	fatty acid binding protein 3, muscle and heart	Hs. 49881. 1	gb:AI041520_RC		1	3. 7	314	2. 50	1. 25	0. 33
672	Heart	214952_at	neural cell adhesion molecule 1	Hs. 167988. 4	gb:BF348061		11	3. 7	293	2. 47	0. 90	0. 42
673	Heart	207501_s_at	fibroblast growth factor 12B	Hs. 326401. 0	gb:NM_004113. 2	FGF12B	3	3. 6	169	2. 23	1. 52	0. 20
674	Heart	214961_at	KIAA0774 protein	Hs. 22201. 0	gb:AI818409_RC		13	3. 4	225	2. 35	0. 71	0. 49
675	Heart	217066_s_at	dystrophia myotonica-protein kinase	Hs. 898. 3	gb:M87313. 1		NA	3. 4	131	2. 12	1. 01	0. 33
676	Heart	203441_s_at	cadherin 2, type 1, N-cadherin (neuronal)	Hs. 161. 0	gb:NM_001792. 1	CDH2	18	3. 2	254	2. 41	1. 57	0. 26
677	Heart	204324_s_at	type II Golgi membrane protein	Hs. 143600. 0	gb:NM_014498. 1	GPP130	3	3. 1	111	2. 04	1. 06	0. 31
678	Heart	217623_at	ESTs	Hs. 130465. 0	gb:BF114815_RC		16	3	120	2. 08	0. 98	0. 36
679	Heart	210121_at	UDP-Gal:betaGlcNAc beta 1, 3-galactosyltransfera	Hs. 181353. 0	gb:AF288390. 1		1	3	71	1. 85	0. 15	0. 56
680	Liver	206753_at	microsomal NAD+-dependent retinol dehydrogenase	Hs. 134958. 0	gb:AF086735. 1		12	20. 3	754	2. 88	0. 94	0. 10
681	Liver	206287_s_at	inter-alpha (globulin) inhibitor H4 (plasma Ka	Hs. 76415. 0	gb:NM_002218. 1	ITIH4	3	17. 3	2809	3. 45	0. 80	0. 15
682	Liver	210049_at	serine (or cysteine) proteinase inhibitor, cla	Hs. 75599. 0	gb:D29832. 1		1	17. 1	3789	3. 58	0. 66	0. 17
683	Liver	204987_at	inter-alpha (globulin) inhibitor, H2 polypeptic	Hs. 75285. 0	gb:NM_002216. 1	ITIH2	10	16. 4	4626	3. 67	1. 84	0. 11
684	Liver	207874_s_at	complement factor H-related 4	Hs. 194776. 0	gb:NM_006684. 1	FHR-4	1	15. 4	1512	3. 18	0. 74	0. 16

Fig. 59

	A	B	C	D	E	F	G	H	I	J	K	L
685	Liver	210327_s_at	alanine-glyoxylate aminotransferase (oxalosis I	Hs. 144567. 0	gb:D13368. 1	AGXT		14. 5	3300	3. 52	0. 32	0. 22
686	Liver	206226_at	histidine-rich glycoprotein	Hs. 1498. 0	gb:NM_000412. 2	HRG	14	14	4129	3. 62	1. 73	0. 13
687	Liver	209976_s_at	cytochrome P450, subfamily IIE (ethanol-inducit	Hs. 75183. 0	gb:AF182276. 1	CYP2E1	10	13. 3	4563	3. 66	0. 49	0. 24
688	Liver	206738_at	apolipoprotein C-IV	Hs. 110675. 0	gb:NM_001646. 1	APOC4	19	13. 2	935	2. 97	1. 84	0. 09
689	Liver	206305_s_at	complement component 8, alpha polypeptide	Hs. 93210. 0	gb:NM_000562. 1	C8A	1	13. 1	1326	3. 12	0. 92	0. 17
690	Liver	208088_s_at			gb:NM_030787. 1	FHR5		12. 4	429	2. 63	0. 05	0. 21
691	Liver	1431_at						12. 3	3940	3. 60	1. 39	0. 18
692	Liver	206910_x_at	H factor (complement)-like 3	Hs. 154224. 0	gb:NM_005666. 1	HFL3	1	12. 2	3416	3. 53	1. 73	0. 15
693	Liver	205754_at	coagulation factor II (thrombin)	Hs. 76530. 0	gb:NM_000506. 2	F2		11. 3	3122	3. 49	1. 13	0. 21
694	Liver	39763_at						11. 2	5417	3. 73	1. 44	0. 21
695	Liver	204998_s_at	activating transcription factor 5	Hs. 9754. 0	gb:NM_012068. 2	ATF5	19	11. 2	1812	3. 26	1. 72	0. 14
696	Liver	206130_s_at	asialoglycoprotein receptor 2	Hs. 1259. 0	gb:NM_001181. 1	ASGR2	17	11. 2	1574	3. 20	0. 83	0. 21
697	Liver	206840_at	afamin	Hs. 531. 0	gb:NM_001133. 1	AFM	4	10. 5	1275	3. 11	1. 62	0. 14
698	Liver	206913_at	bile acid Coenzyme A: amino acid N-acyltransfe	Hs. 159440. 0	gb:NM_001701. 1	BAAT	9	10. 2	379	2. 58	0. 61	0. 19
699	Liver	205943_at	tryptophan 2, 3-dioxygenase	Hs. 183671. 0	gb:NM_005651. 1	TDO2	4	10	2734	3. 44	1. 66	0. 18
700	Liver	206354_at	solute carrier family 21 (organic anion transpo	Hs. 274981. 0	gb:NM_019844. 1	SLC21A8	12	9. 5	469	2. 67	0. 77	0. 20
701	Liver	210929_s_at	Homo sapiens clone FLB5539 PRO1454 mRNA, comple	Hs. 323288. 0	gb:AF130057. 1		12	9. 4	4389	3. 64	0. 57	0. 33
702	Liver	204999_s_at	activating transcription factor 5	Hs. 9754. 0	gb:BC005174. 1		19	9. 1	303	2. 48	1. 41	0. 12
703	Liver	207097_s_at	solute carrier family 17 (sodium phosphate), me	Hs. 19710. 0	gb:NM_005835. 1	SLC17A2	6	9. 1	425	2. 63	0. 55	0. 23
704	Liver	214320_x_at	cytochrome P450, subfamily IIA (phenobarbital-i	Hs. 183584. 2	gb:T67741_RC		5	8. 9	1696	3. 23	1. 80	0. 16
705	Liver	206350_at	amyloid P component, serum	Hs. 1957. 0	gb:NM_001639. 1	APCS	1	8. 7	5377	3. 73	1. 49	0. 26
706	Liver	207300_s_at	coagulation factor VII (serum prothrombin conve	Hs. 36989. 1	gb:NM_000131. 2	F7	13	8. 6	596	2. 78	1. 51	0. 15
707	Liver	214465_at	orosomucoid 2	Hs. 278388. 0	gb:NM_000608. 1	ORM2	9	8. 4	4467	3. 65	0. 87	0. 33
708	Liver	205755_at	pre-alpha (globulin) inhibitor, H3 polypeptide	Hs. 76716. 0	gb:NM_002217. 1	ITIH3	3	8. 3	3921	3. 59	1. 50	0. 25
709	Liver	206606_at	lipase, hepatic	Hs. 9994. 0	gb:NM_000236. 1	LIPC	15	8. 3	331	2. 52	0. 58	0. 23
710	Liver	209975_at	cytochrome P450, subfamily IIE (ethanol-inducit	Hs. 75183. 0	gb:AF182276. 1	CYP2E1	10	8. 3	5914	3. 77	1. 58	0. 26
711	Liver	210521_s_at	fetuin B	Hs. 81073. 0	gb:AB017551. 1	16G2	3	8. 1	180	2. 26	0. 40	0. 23
712	Liver	213632_at	dihydroorotate dehydrogenase	Hs. 94925. 0	gb:M94065. 1		16	8. 1	417	2. 62	1. 67	0. 12
713	Liver	206727_at	complement component 9	Hs. 1290. 0	gb:K02766. 1	C9	5	8	1502	3. 18	0. 87	0. 29
714	Liver	205477_s_at	alpha-1-microglobulinbikunin precursor	Hs. 76177. 0	gb:NM_001633. 1	AMBP	9	7. 9	6511	3. 81	0. 59	0. 41
715	Liver	207409_at	leukocyte cell-derived chemotaxin 2	Hs. 167877. 0	gb:NM_002302. 1	LECT2	5	7. 9	1699	3. 23	0. 91	0. 29
716	Liver	205813_s_at	methionine adenosyltransferase I, alpha	Hs. 323715. 0	gb:NM_000429. 1	MAT1A	10	7. 8	2269	3. 36	1. 05	0. 30
717	Liver	207218_at	coagulation factor IX (plasma thromboplastic co	Hs. 1330. 0	gb:NM_000133. 2	F9	X	7. 8	3218	3. 51	1. 29	0. 29
718	Liver	204551_s_at	alpha-2-HS-glycoprotein	Hs. 324746. 0	gb:NM_001622. 1	AHSG	3	7. 7	3271	3. 51	0. 87	0. 34
719	Liver	207102_at	aldo-keto reductase family 1, member D1 (delta	Hs. 201667. 0	gb:NM_005989. 1	AKR1D1	7	7. 7	918	2. 96	1. 15	0. 23
720	Liver	205351_at	gamma-glutamyl carboxylase	Hs. 77719. 0	gb:NM_000821. 1	GGCX	2	7. 6	432	2. 64	1. 81	0. 11
721	Liver	207820_at	alcohol dehydrogenase 1 (class I), alpha polype	Hs. 73843. 0	gb:NM_000667. 2	ADH1	4	7. 6	1583	3. 20	1. 35	0. 24
722	Liver	210324_at	complement component 8, gamma polypeptide	Hs. 1285. 0	gb:M17263. 1	C8G	9	7. 5	552	2. 74	0. 26	0. 33

Fig. 60

	A	B	C	D	E	F	G	H	I	J	K	L
723	Liver	219466_s_at	apolipoprotein A-II	Hs. 237658. 0	gb:NM_001643. 1	APOA2		7. 4	5260	3. 72	0. 73	0. 40
724	Liver	31835_at						7. 3	1153	3. 06	1. 32	0. 24
725	Liver	205500_at	complement component 5	Hs. 1281. 0	gb:NM_001735. 1	C5	9	7. 3	2017	3. 30	1. 49	0. 25
726	Liver	206651_s_at	carboxypeptidase B-like protein	Hs. 274495. 0	gb:NM_016413. 1	LOC51745	13	7. 3	3699	3. 57	1. 36	0. 30
727	Liver	207027_at	HGF activator	Hs. 104. 0	gb:NM_001528. 1	HGFAC	4	7. 3	744	2. 87	0. 87	0. 27
728	Liver	205216_s_at	apolipoprotein H (beta-2-glycoprotein I)	Hs. 1252. 0	gb:NM_000042. 1	APOH	17	7. 2	9119	3. 96	1. 25	0. 38
729	Liver	204428_s_at	lecithin-cholesterol acyltransferase	Hs. 325507. 0	gb:NM_000229. 1	LCAT	16	7. 1	267	2. 43	0. 56	0. 26
730	Liver	210013_at	hemopexin	Hs. 1504. 0	gb:BC005395. 1		11	7. 1	6157	3. 79	1. 45	0. 33
731	Liver	210888_s_at	inter-alpha (globulin) inhibitor, H1 polypeptic	Hs. 2777. 1	gb:AF116713. 1_RC		3	7. 1	2743	3. 44	1. 05	0. 34
732	Liver	206755_at	cytochrome P450, subfamily IIB (phenobarbital-i	Hs. 1360. 0	gb:NM_000767. 2	CYP2B6	19	6. 9	628	2. 80	1. 07	0. 25
733	Liver	37201_at						6. 8	493	2. 69	1. 88	0. 12
734	Liver	204561_x_at	apolipoprotein C-II	Hs. 75615. 0	gb:NM_000483. 2	APOC2	19	6. 8	6507	3. 81	1. 38	0. 36
735	Liver	217133_x_at	cytochrome P450, subfamily IIB (phenobarbital-i	Hs. 1360. 1	gb:X06399. 1		19	6. 8	142	2. 15	1. 37	0. 12
736	Liver	220626_at	protein Z-dependent protease inhibitor precurso	Hs. 118620. 0	gb:NM_016186. 1	LOC51156	14	6. 8	748	2. 87	0. 41	0. 36
737	Liver	205719_s_at	phenylalanine hydroxylase	Hs. 1870. 0	gb:NM_000277. 1	PAH	12	6. 7	1467	3. 17	1. 18	0. 30
738	Liver	206256_at	carboxypeptidase N, polypeptide 1, 50kD	Hs. 2246. 0	gb:NM_001308. 1	CPN1	10	6. 7	411	2. 61	1. 78	0. 13
739	Liver	206259_at	protein C (inactivator of coagulation factors V	Hs. 2351. 0	gb:NM_000312. 1	PROC	2	6. 7	345	2. 54	0. 58	0. 29
740	Liver	208209_s_at	complement component 4-binding protein, beta	Hs. 99886. 0	gb:NM_000716. 1	C4BPB		6. 7	970	2. 99	0. 92	0. 31
741	Liver	206979_at	complement component 8, beta polypeptide	Hs. 38069. 0	gb:NM_000066. 1	C8B	1	6. 6	1999	3. 30	1. 61	0. 25
742	Liver	207330_at	pregnancy-zone protein	Hs. 74094. 0	gb:NM_002864. 1	PZP	12	6. 6	527	2. 72	1. 93	0. 12
743	Liver	217558_at	ESTs, Moderately similar to 1506290A cytochrome	Hs. 282639. 0	gb:BE971373		10	6. 5	482	2. 68	1. 57	0. 17
744	Liver	207256_at	mannose-binding lectin (protein C) 2, soluble (	Hs. 2314. 0	gb:NM_000242. 1	MBL2	10	6. 4	934	2. 97	0. 60	0. 37
745	Liver	208034_s_at	protein Z, vitamin K-dependent plasma glycoprot	Hs. 1011. 0	gb:NM_003891. 1	PROZ	13	6. 4	323	2. 51	1. 71	0. 13
746	Liver	206345_s_at	paraoxonase 1	Hs. 1898. 0	gb:NM_000446. 1	PON1	7	6. 3	1844	3. 27	1. 44	0. 29
747	Liver	207498_s_at	cytochrome P450, subfamily IID (debrisoquine, s	Hs. 301086. 0	gb:NM_000106. 1	CYP2D6	22	6. 3	1894	3. 28	1. 49	0. 28
748	Liver	221605_s_at	L-pipecolic acid oxidase	Hs. 271167. 1	gb:AF136970. 1		17	6. 3	654	2. 82	1. 70	0. 18
749	Liver	205871_at	plasminogen-like	Hs. 262869. 0	gb:BC005379. 1		2	6. 1	931	2. 97	1. 12	0. 30
750	Liver	208147_s_at			gb:NM_030878. 1	CYP2C8	10	6. 1	1757	3. 24	1. 14	0. 35
751	Liver	220224_at	hydroxyacid oxidase (glycolate oxidase) 1	Hs. 193640. 0	gb:NM_017545. 2	HAO1	20	6. 1	883	2. 95	1. 36	0. 26
752	Liver	206386_at	serine (or cysteine) proteinase inhibitor, clad	Hs. 76838. 0	gb:NM_000354. 1	SERPINA7	X	6	1201	3. 08	1. 26	0. 30
753	Liver	210798_x_at	mannan-binding lectin serine protease 2	Hs. 119983. 1	gb:AB008047. 1	sMAP	1	6	336	2. 53	0. 95	0. 26
754	Liver	206505_at	UDP glycosyltransferase 2 family, polypeptide B	Hs. 89691. 0	gb:NM_021139. 1	UGT2B4	4	5. 9	889	2. 95	0. 03	0. 49
755	Liver	219465_at	apolipoprotein A-II	Hs. 237658. 0	gb:NM_001643. 1	APOA2		5. 9	7081	3. 85	0. 90	0. 50
756	Liver	207584_at	lipoprotein, Lp(a)	Hs. 119520. 0	gb:NM_005577. 1	LPA	6	5. 8	212	2. 33	0. 61	0. 29
757	Liver	207185_at	solute carrier family 10 (sodiumbile acid cotra	Hs. 952. 0	gb:NM_003049. 1	SLC10A1	14	5. 7	1296	3. 11	1. 90	0. 21
758	Liver	207201_s_at	solute carrier family 22 (organic cation transp	Hs. 117367. 0	gb:NM_003057. 1	SLC22A1	6	5. 7	1227	3. 09	0. 74	0. 41
759	Liver	210326_at	alanine-glyoxylate aminotransferase (oxalosis I	Hs. 144567. 0	gb:D13368. 1	AGXT		5. 7	900	2. 95	1. 58	0. 24
760	Liver	219559_at	hypothetical protein FLJ23412	Hs. 34487. 0	gb:NM_022082. 1	FLJ23412	20	5. 7	427	2. 63	1. 91	0. 13

Fig. 61

	A	B	C	D	E	F	G	H	I	J	K	L
761	Liver	203775_at	solute carrier family 25, member 13 (citrin)	Hs. 9599. 0	gb:NM_014251. 1	SLC25A13	7	5. 6	300	2. 48	1. 73	0. 13
762	Liver	207995_s_at	CD209 antigen-like	Hs. 23759. 0	gb:NM_014257. 1	CD209L	19	5. 6	342	2. 53	1. 51	0. 18
763	Liver	203500_at	glutaryl-Coenzyme A dehydrogenase	Hs. 184141. 1	gb:NM_000159. 1	GCDH	19	5. 5	380	2. 58	1. 76	0. 15
764	Liver	205306_x_at	kynurenine 3-monooxygenase (kynurenine 3-hydrox	Hs. 107318. 0	gb:AI074145_RC		1	5. 5	484	2. 69	1. 69	0. 18
765	Liver	210558_at	aldo-keto reductase family 1, member C4 (chlord	Hs. 177687. 0	gb:AB045829. 1		10	5. 5	348	2. 54	0. 84	0. 31
766	Liver	40665_at						5. 4	951	2. 98	1. 17	0. 33
767	Liver	220801_s_at	hydroxyacid oxidase 2 (long chain)	Hs. 236545. 0	gb:NM_016527. 1	HAO2	1	5. 4	440	2. 64	1. 83	0. 15
768	Liver	213631_x_at	dihydroorotate dehydrogenase	Hs. 94925. 0	gb:BE789211		16	5. 3	259	2. 41	1. 71	0. 13
769	Liver	207041_at	mannan-binding lectin serine protease 2	Hs. 119983. 0	gb:NM_006610. 1	MASP2	1	5. 2	358	2. 55	1. 72	0. 16
770	Liver	204880_at	O-6-methylguanine-DNA methyltransferase	Hs. 1384. 0	gb:NM_002412. 1	MGMT	10	5. 1	291	2. 46	1. 81	0. 13
771	Liver	206867_at	glucokinase (hexokinase 4) regulatory protein	Hs. 89771. 0	gb:NM_001486. 1	GCKR	2	5. 1	340	2. 53	1. 72	0. 16
772	Liver	207773_x_at	cytochrome P450 polypeptide 43	Hs. 306220. 0	gb:NM_022820. 2	CYP3A43	7	5. 1	255	2. 41	0. 73	0. 33
773	Liver	210366_at	solute carrier family 21 (organic anion transpo	Hs. 137425. 0	gb:AB026257. 1	OATP-C	12	5. 1	885	2. 95	0. 79	0. 42
774	Liver	211138_s_at	kynurenine 3-monooxygenase (kynurenine 3-hydrox	Hs. 107318. 1	gb:BC005297. 1		1	5. 1	615	2. 79	1. 77	0. 20
775	Liver	219733_s_at	very long-chain acyl-CoA synthetase homolog 2	Hs. 111401. 0	gb:NM_012254. 1	VLCS-H2	5	5. 1	895	2. 95	1. 28	0. 33
776	Liver	221159_at	clone FLB1727	Hs. 283773. 0	gb:NM_016414. 1	LOC51215		5. 1	291	2. 46	1. 60	0. 17
777	Liver	204140_at	tyrosylprotein sulfotransferase 1	Hs. 17279. 0	gb:NM_003596. 1	TPST1	7	5	249	2. 40	1. 79	0. 12
778	Liver	205972_at	transporter protein; system N1 Na+ and H+-coupl	Hs. 76460. 0	gb:NM_006841. 1	G17	3	5	326	2. 51	1. 43	0. 21
779	Liver	207007_at	nuclear receptor subfamily 1, group I, member 3	Hs. 83623. 0	gb:NM_005122. 1	NR1I3		5	197	2. 29	1. 54	0. 15
780	Liver	205774_at	coagulation factor XII (Hageman factor)	Hs. 1321. 0	gb:NM_000505. 2	F12		4. 9	2024	3. 31	1. 30	0. 41
781	Liver	206153_at	cytochrome P450, subfamily IVF, polypeptide 11	Hs. 187393. 0	gb:NM_021187. 1	CYP4F11	19	4. 9	75	1. 88	0. 50	0. 28
782	Liver	216223_at	carboxypeptidase N, polypeptide 2, 83kD	Hs. 73858. 0	gb:J05158. 1	ACBP	3	4. 9	448	2. 65	1. 36	0. 27
783	Liver	217512_at	ESTs, Highly similar to 1312232A kininogen L, hi	Hs. 289056. 0	gb:BG398937		3	4. 9	811	2. 91	1. 33	0. 32
784	Liver	220507_s_at	beta-ureidopropionase	Hs. 126926. 0	gb:NM_016327. 1	LOC51733	22	4. 9	733	2. 86	0. 79	0. 42
785	Liver	221142_s_at	peroxisomal trans 2-enoyl CoA reductase; putati	Hs. 281680. 0	gb:NM_018441. 1	HSA250303	2	4. 9	212	2. 33	1. 23	0. 23
786	Liver	89977_at						4. 8	506	2. 70	1. 82	0. 19
787	Liver	202959_at	methylmalonyl Coenzyme A mutase	Hs. 155212. 0	gb:AI433712_RC			4. 7	401	2. 60	1. 84	0. 16
788	Liver	207544_s_at	alcohol dehydrogenase 6 (class V)	Hs. 76800. 0	gb:NM_000672. 2	ADH6	4	4. 7	327	2. 51	1. 11	0. 30
789	Liver	57539_at						4. 6	188	2. 27	1. 64	0. 14
790	Liver	213596_at	Homo sapiens mRNA; cDNA DKFZp586A181 (from clor	Hs. 321247. 0	gb:AL050391. 1	DKFZp586A1	11	4. 6	183	2. 26	1. 68	0. 12
791	Liver	214282_at	ceruloplasmin (ferroxidase)	Hs. 296634. 1	gb:AA191647_RC		3	4. 6	123	2. 09	0. 83	0. 28
792	Liver	214621_at	glycogen synthase 2 (liver)	Hs. 82614. 0	gb:S70004. 1	glycogen s	12	4. 6	391	2. 59	0. 93	0. 36
793	Liver	215206_at	Homo sapiens cDNA: FLJ21490 fis, clone COL05464	Hs. 288700. 0	gb:AK025143. 1		8	4. 6	424	2. 63	1. 72	0. 20
794	Liver	206325_at	serine (or cysteine) proteinase inhibitor, clad	Hs. 1305. 0	gb:NM_001756. 2	SERPINA6	14	4. 5	1002	3. 00	1. 41	0. 36
795	Liver	206743_s_at	asialoglycoprotein receptor 1	Hs. 12056. 0	gb:NM_001671. 1	ASGR1	17	4. 5	743	2. 87	0. 99	0. 42
796	Liver	207810_at	coagulation factor XIII, B polypeptide	Hs. 68601. 0	gb:NM_001994. 1	F13B	1	4. 5	258	2. 41	0. 98	0. 32
797	Liver	207907_at	tumor necrosis factor (ligand) superfamily, mem	Hs. 129708. 0	gb:NM_003807. 1	TNFSF14	19	4. 5	213	2. 33	0. 65	0. 37
798	Liver	210063_at	sarcosine dehydrogenase	Hs. 198003. 3	gb:AF162428. 1		9	4. 5	193	2. 28	1. 20	0. 24

Fig. 62

	A	B	C	D	E	F	G	H	I	J	K	L
799	Liver	220604_x_at	formiminotransferase cyclodeaminase	Hs. 36218. 0	gb:NM_006657. 1	FTCD	21	4. 5	687	2. 84	1. 46	0. 31
800	Liver	220786_s_at	hypothetical protein FLJ10191	Hs. 165655. 0	gb:NM_018018. 1	FLJ10191	12	4. 5	407	2. 61	0. 78	0. 41
801	Liver	221662_s_at	solute carrier family 22 (organic anion transp	Hs. 251395. 1	gb:AF210455. 1		6	4. 5	477	2. 68	1. 55	0. 25
802	Liver	206085_s_at	cystathionase (cystathionine gamma-lyase)	Hs. 19904. 0	gb:NM_001902. 1	CTH	1	4. 4	264	2. 42	1. 26	0. 26
803	Liver	219803_at	angiopoietin-like 3	Hs. 209153. 0	gb:NM_014495. 1	ANGPTL3	1	4. 4	997	3. 00	0. 59	0. 55
804	Liver	220061_at	hypothetical protein FLJ20581	Hs. 122939. 0	gb:NM_017888. 1	FLJ20581		4. 4	450	2. 65	0. 98	0. 38
805	Liver	220496_at	C-type lectin-like receptor-2	Hs. 114231. 0	gb:NM_016509. 1	LOC51266	12	4. 4	409	2. 61	1. 53	0. 25
806	Liver	206938_at	steroid-5-alpha-reductase, alpha polypeptide 2	Hs. 1989. 0	gb:NM_000348. 1	SRD5A2	2	4. 3	177	2. 25	1. 47	0. 18
807	Liver	217319_x_at	cytochrome P450, subfamily IVA, polypeptide 11	Hs. 1645. 2	gb:AL135960			4. 3	586	2. 77	1. 13	0. 38
808	Liver	217521_at	ESTs	Hs. 276590. 0	gb:N54942_RC		12	4. 3	488	2. 69	1. 44	0. 29
809	Liver	205923_at	reelin	Hs. 12246. 0	gb:NM_005045. 1	RELN	7	4. 2	328	2. 52	1. 68	0. 20
810	Liver	205939_at	cytochrome P450, subfamily IIIA, polypeptide 7	Hs. 172323. 0	gb:NM_000765. 2	CYP3A7	7	4. 2	977	2. 99	1. 36	0. 39
811	Liver	215078_at	Homo sapiens mRNA; cDNA DKFZp564M2422 (from cl	Hs. 306320. 0	gb:AL050388. 1	DKFZp564M2	6	4. 2	336	2. 53	1. 28	0. 30
812	Liver	205984_at	corticotropin releasing hormone-binding protei	Hs. 115617. 0	gb:NM_001882. 2	CRHBP	5	4. 1	539	2. 73	1. 48	0. 31
813	Liver	206610_s_at	coagulation factor XI (plasma thromboplastin a	Hs. 1430. 1	gb:NM_000128. 2	F11	4	4. 1	266	2. 43	1. 47	0. 23
814	Liver	207883_s_at	transferrin receptor 2	Hs. 63758. 0	gb:NM_003227. 1	TFR2	7	4	301	2. 48	0. 92	0. 39
815	Liver	214478_at	secreted phosphoprotein 2, 24kD	Hs. 12230. 0	gb:NM_006944. 1	SPP2	2	4	559	2. 75	1. 67	0. 27
816	Liver	216687_x_at	UDP glycosyltransferase 2 family, polypeptide	Hs. 150207. 1	gb:U06641. 1		4	4	574	2. 76	1. 09	0. 41
817	Liver	217127_at	cystathionase (cystathionine gamma-lyase)	Hs. 19904. 1	gb:AL354872		1	4	521	2. 72	1. 43	0. 32
818	Liver	206541_at	kallikrein B, plasma (Fletcher factor) 1	Hs. 1901. 0	gb:NM_000892. 2	KLKB1	4	3. 9	373	2. 57	1. 13	0. 37
819	Liver	207262_at	apolipoprotein F	Hs. 2388. 0	gb:NM_001638. 1	APOF	12	3. 9	358	2. 55	0. 43	0. 54
820	Liver	207819_s_at	ATP-binding cassette, sub-family B (MDRTAP), m	Hs. 73812. 1	gb:NM_000443. 2	ABCB4	7	3. 9	503	2. 70	1. 15	0. 40
821	Liver	222078_at	KIAA1535 protein	Hs. 284171. 0	gb:BF110802_RC			3. 9	126	2. 10	0. 64	0. 37
822	Liver	205620_at	coagulation factor X	Hs. 47913. 0	gb:NM_000504. 2	F10	13	3. 8	217	2. 34	1. 40	0. 25
823	Liver	209614_at	alcohol dehydrogenase 2 (class I), beta polype	Hs. 4. 0	gb:AF153821. 1		4	3. 8	486	2. 69	1. 15	0. 41
824	Liver	214415_at	plasminogen-like	Hs. 262869. 1	gb:N58120_RC		2	3. 8	194	2. 29	1. 03	0. 33
825	Liver	221009_s_at	PPAR(gamma) angiopoietin related protein	Hs. 9613. 0	gb:NM_016109. 1	PGAR	19	3. 8	173	2. 24	1. 24	0. 26
826	Liver	206449_s_at	mannan-binding lectin serine protease 1 (C4C2 a	Hs. 227152. 0	gb:NM_001879. 1	MASP1	3	3. 7	141	2. 15	1. 45	0. 19
827	Liver	206496_at	flavin containing monooxygenase 3	Hs. 80876. 0	gb:NM_006894. 2	FMO3	1	3. 7	1014	3. 01	0. 95	0. 55
828	Liver	207244_x_at	cytochrome P450, subfamily IIA (phenobarbital-	Hs. 183584. 0	gb:NM_000762. 2	CYP2A6	19	3. 7	1001	3. 00	1. 17	0. 49
829	Liver	209916_at	hypothetical protein DKFZp762M115	Hs. 271586. 0	gb:BC002477. 1		10	3. 7	220	2. 34	1. 55	0. 21
830	Liver	211354_s_at	leptin receptor	Hs. 226627. 2	gb:U52913. 1			3. 7	225	2. 35	1. 45	0. 24
831	Liver	211442_x_at	cytochrome P450 polypeptide 43	Hs. 306220. 3	gb:AF280111. 1	CYP3A43	7	3. 7	237	2. 37	0. 85	0. 41
832	Liver	214837_at	albumin	Hs. 184411. 7	gb:M12523		4	3. 7	186	2. 27	0. 93	0. 36
833	Liver	206952_at	glucose-6-phosphatase, catalytic (glycogen sto	Hs. 242. 0	gb:NM_000151. 1	G6PC	17	3. 6	235	2. 37	0. 46	0. 54
834	Liver	207804_s_at	ficolin (collagenfibrinogen domain-containing	Hs. 54517. 3	gb:NM_004108. 1	FCN2		3. 6	253	2. 40	1. 06	0. 37
835	Liver	207609_s_at	cytochrome P450, subfamily I (aromatic compoun	Hs. 1361. 0	gb:AF182274. 1	CYP1A2	15	3. 5	161	2. 21	1. 14	0. 30
836	Liver	217583_at	ESTs	Hs. 297257. 0	gb:BG433489		12	3. 5	90	1. 96	0. 68	0. 36

Fig. 63

	A	B	C	D	E	F	G	H	I	J	K	L
837	Liver	210233_at	interleukin 1 receptor accessory protein	Hs. 173880. 1	gb:AF167343. 1	IL1RAP	3	3. 4	225	2. 35	0. 67	0. 50
838	Liver	210328_at	glycine N-methyltransferase	Hs. 144914. 0	gb:AF101477. 1	GNMT	6	3. 4	297	2. 47	0. 94	0. 45
839	Liver	206957_at	alanine-glyoxylate aminotransferase homolog	Hs. 271366. 0	gb:NM_016236. 1	TLH6		3. 3	88	1. 94	0. 76	0. 36
840	Liver	210587_at	Homo sapiens, Similar to inhibin beta E, clone	Hs. 279497. 0	gb:BC005161. 1		12	3. 3	197	2. 30	1. 01	0. 39
841	Liver	214307_at	homogentisate 1, 2-dioxygenase (homogentisate ox	Hs. 15113. 1	gb:AI478172_RC		3	3. 3	352	2. 55	1. 19	0. 41
842	Liver	214308_s_at	homogentisate 1, 2-dioxygenase (homogentisate ox	Hs. 15113. 1	gb:AI478172_RC		3	3. 3	419	2. 62	1. 21	0. 42
843	Liver	205222_at	enoyl-Coenzyme A, hydratase3-hydroxyacyl Coenzy	Hs. 1531. 0	gb:NM_001966. 1	EHHADH	3	3. 2	228	2. 36	0. 99	0. 43
844	Liver	205307_s_at	kynurenine 3-monooxygenase (kynurenine 3-hydrox	Hs. 107318. 0	gb:NM_003679. 1	KMO	1	3. 2	187	2. 27	1. 05	0. 38
845	Liver	210680_s_at	mannan-binding lectin serine protease 1 (C4C2 a	Hs. 227152. 2	gb:D17525. 1	CRARF	3	3. 2	140	2. 15	1. 55	0. 19
846	Liver	220554_at	solute carrier family 22 (organic anion transpo	Hs. 251395. 0	gb:NM_006672. 1	SLC22A7	6	3. 2	96	1. 98	0. 72	0. 40
847	Liver	204973_at	gap junction protein, beta 1, 32kD (connexin 32	Hs. 2679. 0	gb:NM_000166. 1	GJB1	X	3. 1	285	2. 45	1. 67	0. 26
848	Liver	220528_at	VNN3 protein	Hs. 183656. 0	gb:NM_018399. 1	HSA238982	6	3. 1	143	2. 15	0. 93	0. 40
849	Liver	206155_at	ATP-binding cassette, sub-family C (CFTR/MRP), m	Hs. 193852. 0	gb:NM_000392. 1	ABCC2	10	3	215	2. 33	1. 12	0. 40
850	Liver	207354_at	small inducible cytokine subfamily A (Cys-Cys),	Hs. 10458. 0	gb:NM_004590. 1	SCYA16	17	2. 9	181	2. 26	1. 45	0. 28
851	Liver	211248_s_at	chordin	Hs. 166186. 2	gb:AF283325. 1	CHRD	3	2. 8	223	2. 35	1. 54	0. 29
852	Liver	207406_at	cytochrome P450, subfamily VIIA (cholesterol 7	Hs. 1644. 0	gb:NM_000780. 2	CYP7A1	8	2. 5	64	1. 81	0. 65	0. 46
853	Spleen	220832_at	Toll-like receptor 8	Hs. 272410. 0	gb:NM_016610. 1	LOC51311	X	8. 5	96	1. 98	0. 46	0. 18
854	Spleen	210225_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 105928. 0	gb:AF009635. 1		19	6. 7	267	2. 43	0. 98	0. 21
855	Spleen	210784_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 105928. 2	gb:AF009634. 1		19	6. 3	223	2. 35	0. 94	0. 22
856	Spleen	211135_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 105928. 4	gb:AF009644. 1		19	5. 5	218	2. 34	0. 98	0. 25
857	Spleen	207104_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 204040. 0	gb:NM_006669. 1	LILRB1	19	5. 1	170	2. 23	0. 88	0. 26
858	Spleen	207610_s_at	egf-like module containing, mucin-like, hormone	Hs. 137354. 0	gb:NM_013447. 1	EMR2	19	4. 8	323	2. 51	1. 50	0. 21
859	Spleen	210176_at	toll-like receptor 1	Hs. 2474. 0	gb:AL050262. 1	DKFZp564I0	4	4. 6	147	2. 17	1. 37	0. 17
860	Spleen	206244_at	complement component (3b4b) receptor 1, includi	Hs. 193716. 0	gb:NM_000573. 1	CR1	1	4. 5	117	2. 07	0. 81	0. 28
861	Spleen	208488_s_at	complement component (3b4b) receptor 1, includi	Hs. 193716. 1	gb:NM_000651. 1	CR1	1	4. 4	196	2. 29	1. 55	0. 17
862	Spleen	206336_at	small inducible cytokine subfamily B (Cys-X-Cys	Hs. 164021. 0	gb:NM_002993. 1	SCYB6	4	4. 1	247	2. 39	1. 40	0. 24
863	Spleen	206420_at	immunoglobulin superfamily, member 6	Hs. 135194. 0	gb:NM_005849. 1	IGSF6	16	4. 1	379	2. 58	1. 31	0. 31
864	Spleen	210786_s_at	Friend leukemia virus integration 1	Hs. 108043. 1	gb:M93255. 1	FLI-1; FLI	11	4	237	2. 37	1. 68	0. 17
865	Spleen	211133_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 105928. 1	gb:AF009643. 1		19	3. 9	199	2. 30	1. 36	0. 24
866	Spleen	206467_x_at	tumor necrosis factor receptor superfamily, mem	Hs. 278556. 0	gb:NM_003823. 1	TNFRSF6B	20	3. 6	246	2. 39	1. 79	0. 17
867	Spleen	207008_at	interleukin 8 receptor, beta	Hs. 846. 0	gb:NM_001557. 1	IL8RB	2	3. 6	253	2. 40	0. 72	0. 47
868	Spleen	221060_s_at	toll-like receptor 4	Hs. 159239. 0	gb:NM_003266. 1	TLR4	9	3. 6	275	2. 44	1. 61	0. 23
869	Spleen	211336_x_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 204040. 2	gb:AF009007. 1		19	3. 5	241	2. 38	1. 23	0. 33
870	Spleen	220577_at	hypothetical protein FLJ13373	Hs. 287567. 0	gb:NM_025006. 1	FLJ13373	11	3. 3	256	2. 41	1. 71	0. 21
871	Spleen	210660_at	leukocyte immunoglobulin-like receptor, subfami	Hs. 166156. 1	gb:AF025529. 1	LIR-6	19	3. 1	112	2. 05	0. 83	0. 39
872	Spleen	204860_s_at	baculoviral IAP repeat-containing 1	Hs. 79019. 0	gb:AI817801_RC		5	3	194	2. 29	0. 93	0. 45
873	Spleen	206214_at	phospholipase A2, group VII (platelet-activatin	Hs. 93304. 0	gb:NM_005084. 1	PLA2G7	6	3	247	2. 39	1. 55	0. 28
874	Spleen	209829_at	chromosome 6 open reading frame 32	Hs. 101359. 1	gb:AB002384. 1	KIAA0386	6	3	365	2. 56	1. 55	0. 33

Fig. 64

	A	B	C	D	E	F	G	H	I	J	K	L
875	Spleen	206618_at	interleukin 18 receptor 1	Hs. 159301. 0	gb:NM_003855. 1	IL18R1	2	2. 6	109	2. 04	1. 20	0. 32
876	Kidney	210403_s_at	potassium inwardly-rectifying channel, subfami	Hs. 463. 1	gb:U12543. 1		11	12	409	2. 61	0. 92	0. 14
877	Kidney	221298_s_at	solute carrier family 22 (organic anion transp	Hs. 266223. 0	gb:NM_004254. 1	SLC22A8	11	9. 1	275	2. 44	0. 90	0. 17
878	Kidney	220424_at	NPHS2 gene (podocin)	Hs. 192657. 0	gb:NM_014625. 1	NPHS2	1	7. 4	926	2. 97	1. 05	0. 26
879	Kidney	205244_s_at	sodium-dependent high-affinity dicarboxylate t	Hs. 102867. 0	gb:NM_022829. 1	NADC3	20	7. 2	1207	3. 08	0. 92	0. 30
880	Kidney	206716_at	uromodulin (uromucoid, Tamm-Horsfall glycoprot	Hs. 1137. 0	gb:NM_003361. 1	UMOD		6. 9	5174	3. 71	1. 19	0. 37
881	Kidney	205676_at	cytochrome P450, subfamily XXVIIB (25-hydroxyv	Hs. 199270. 0	gb:NM_000785. 1	CYP27B1	12	6. 8	965	2. 98	1. 59	0. 21
882	Kidney	206872_at	solute carrier family 17 (sodium phosphate), m	Hs. 100001. 0	gb:NM_005074. 1	SLC17A1	6	6. 3	419	2. 62	1. 78	0. 13
883	Kidney	210402_at	potassium inwardly-rectifying channel, subfami	Hs. 463. 1	gb:U03884. 1	ROMK1	11	6. 1	518	2. 71	0. 89	0. 30
884	Kidney	220281_at	solute carrier family 12 (sodiumpotassiumchlor	Hs. 123116. 0	gb:AI632015_RC		15	5. 8	727	2. 86	0. 61	0. 39
885	Kidney	205626_s_at	calbindin 1, (28kD)	Hs. 65425. 0	gb:NM_004929. 2	CALB1	8	5. 7	394	2. 60	1. 29	0. 23
886	Kidney	216599_x_at	solute carrier family 22 (organic anion transp	Hs. 23965. 2	gb:AJ271205. 1	OAT1-4		5. 6	163	2. 21	0. 53	0. 30
887	Kidney	205243_at	sodium-dependent high-affinity dicarboxylate t	Hs. 102867. 0	gb:AW466889_RC		20	5. 3	920	2. 96	1. 72	0. 24
888	Kidney	206517_at	cadherin 16, KSP-cadherin	Hs. 115418. 0	gb:NM_004062. 1	CDH16	16	5. 2	1258	3. 10	0. 83	0. 44
889	Kidney	206963_s_at	putative N-acetyltransferase Camello 2	Hs. 278983. 0	gb:NM_016347. 1	CML2	2	4. 7	680	2. 83	1. 20	0. 35
890	Kidney	209425_at	alpha-methylacyl-CoA racemase	Hs. 128749. 0	gb:AA888589_RC		5	4. 3	279	2. 45	1. 42	0. 24
891	Kidney	210199_at	crystallin, alpha A	Hs. 184085. 0	gb:U66584. 1	CRYAA	21	4. 3	443	2. 65	1. 50	0. 27
892	Kidney	205532_s_at	cadherin 6, type 2, K-cadherin (fetal kidney)	Hs. 32963. 0	gb:AU151483_RC		5	4. 2	149	2. 17	1. 18	0. 24
893	Kidney	209424_s_at	alpha-methylacyl-CoA racemase	Hs. 128749. 0	gb:AI796120_RC		5	4. 2	502	2. 70	1. 46	0. 29
894	Kidney	218701_at	CGI-83 protein	Hs. 118554. 0	gb:NM_016027. 1	LOC51110	8	3. 9	314	2. 50	1. 76	0. 19
895	Kidney	203158_s_at	glutaminase	Hs. 239189. 0	gb:AF097493. 1		2	3. 7	220	2. 34	1. 56	0. 21
896	Kidney	210602_s_at	cadherin 6, type 2, K-cadherin (fetal kidney)	Hs. 32963. 1	gb:BC000019. 1		5	3. 5	225	2. 35	1. 02	0. 38
897	Kidney	216230_x_at	sphingomyelin phosphodiesterase 1, acid lysoso	Hs. 77813. 2	gb:M59917			3. 5	91	1. 96	1. 27	0. 20
898	Kidney	210374_x_at	prostaglandin E receptor 3 (subtype EP3)	Hs. 170917. 5	gb:D38300. 1		1	3. 4	347	2. 54	1. 54	0. 29
899	Kidney	203946_s_at	arginase, type II	Hs. 172851. 0	gb:U75667. 1		14	3. 3	406	2. 61	1. 52	0. 33
900	Kidney	210343_s_at	solute carrier family 22 (organic anion transp	Hs. 23965. 1	gb:AF124373. 1	SLC22A6	11	3. 3	129	2. 11	1. 04	0. 32
901	Kidney	219271_at	hypothetical protein FLJ12691	Hs. 15830. 0	gb:NM_024572. 1	FLJ12691	2	3. 3	220	2. 34	1. 27	0. 33
902	Kidney	221590_s_at	methylmalonate-semialdehyde dehydrogenase	Hs. 293970. 1	gb:AF130089. 1_RC		14	3. 3	276	2. 44	1. 70	0. 23
903	Kidney	222281_s_at	ESTs	Hs. 293472. 0	gb:AW517716_RC		4	3. 2	235	2. 37	1. 30	0. 33
904	Kidney	206286_s_at	teratocarcinoma-derived growth factor 1	Hs. 75561. 0	gb:NM_003212. 1	TDGF1	3	3. 1	242	2. 38	1. 10	0. 42
905	Kidney	219773_at	NADPH oxidase 4	Hs. 93847. 0	gb:NM_016931. 1	NOX4	11	3. 1	216	2. 33	1. 37	0. 31
906	Kidney	206228_at	paired box gene 2	Hs. 155644. 0	gb:AW769732_RC		10	3	179	2. 25	1. 21	0. 35
907	Kidney	220503_at	solute carrier family 13 (sodiumsulphate sympo	Hs. 124664. 0	gb:AF260824. 1			3	68	1. 83	0. 32	0. 50
908	Adrenal glan	214622_at	cytochrome P450, subfamily XXIA (steroid 21-hy	Hs. 278430. 0	gb:M17252. 1	CYP21	6	20. 1	3968	3. 60	0. 89	0. 13
909	Adrenal glan	214610_at	cytochrome P450, subfamily XIB (steroid 11-beta	Hs. 301118. 0	gb:AV702430		8	14. 4	4702	3. 67	1. 69	0. 14
910	Adrenal glan	204501_at	nephroblastoma overexpressed gene	Hs. 235935. 0	gb:NM_002514. 1	NOV	8	9	1949	3. 29	1. 87	0. 16
911	Adrenal glan	206294_at	hydroxy-delta-5-steroid dehydrogenase, 3 beta-	Hs. 825. 0	gb:NM_000198. 1	HSD3B2	1	7. 1	5642	3. 75	1. 26	0. 35
912	Adrenal glan	215834_x_at	Homo sapiens scavenger receptor class B type I	Hs. 283965. 0	gb:AV708130		12	6. 7	448	2. 65	0. 92	0. 26

Fig. 65

	A	B	C	D	E	F	G	H	I	J	K	L
913	Adrenal glan	214630_at	cytochrome P450, subfamily XIB (steroid 11-beta	Hs. 184927. 0	gb:X54741. 1	human CYPX	8	6. 6	384	2. 58	0. 82	0. 27
914	Adrenal glan	208291_s_at	tyrosine hydroxylase	Hs. 178237. 0	gb:NM_000360. 1	TH		6. 5	373	2. 57	0. 63	0. 30
915	Adrenal glan	214321_at	nephroblastoma overexpressed gene	Hs. 235935. 1	gb:BF440025_RC		8	6. 4	2316	3. 36	1. 57	0. 28
916	Adrenal glan	201633_s_at	cytochrome b5 outer mitochondrial membrane prec	Hs. 79103. 0	gb:AW235051_RC		16	5. 6	324	2. 51	1. 76	0. 13
917	Adrenal glan	206793_at	phenylethanolamine N-methyltransferase	Hs. 1892. 0	gb:NM_002686. 1	PNMT	17	5. 6	355	2. 55	1. 25	0. 23
918	Adrenal glan	201634_s_at	cytochrome b5 outer mitochondrial membrane prec	Hs. 79103. 0	gb:NM_030579. 1	CYB5-M	16	5. 3	204	2. 31	1. 60	0. 13
919	Adrenal glan	209279_s_at	NAD(P) dependent steroid dehydrogenase-like; H1	Hs. 57698. 0	gb:BC000245. 1		X	5. 1	181	2. 26	1. 69	0. 11
920	Adrenal glan	206450_at	dopamine beta-hydroxylase (dopamine beta-monoox	Hs. 2301. 0	gb:NM_000787. 1	DBH	9	5	191	2. 28	0. 96	0. 26
921	Adrenal glan	208568_at	melanocortin 2 receptor (adrenocorticotropic ho	Hs. 248144. 0	gb:NM_000529. 1	MC2R	18	4. 7	397	2. 60	1. 74	0. 18
922	Adrenal glan	202067_s_at	low density lipoprotein receptor (familial hype	Hs. 213289. 0	gb:AI861942_RC		19	4. 6	414	2. 62	0. 87	0. 38
923	Adrenal glan	206258_at	alpha2, 8-sialyltransferase	Hs. 22985. 0	gb:NM_013305. 1	HSU91641	18	4. 4	69	1. 84	0. 63	0. 27
924	Adrenal glan	208404_x_at	potassium inwardly-rectifying channel, subfamil	Hs. 193044. 0	gb:NM_000890. 2	KCNJ5	11	4. 4	225	2. 35	1. 34	0. 23
925	Adrenal glan	207813_s_at	ferredoxin reductase	Hs. 69745. 1	gb:NM_004110. 2	FDXR	17	4. 3	1321	3. 12	1. 38	0. 41
926	Adrenal glan	211572_s_at	solute carrier family 23 (nucleobase transporte	Hs. 82042. 1	gb:AF092511. 1	SLC23A1	20	3. 9	147	2. 17	1. 57	0. 15
927	Adrenal glan	215665_at	Human DNA sequence from clone RP5-871G17 on ch	Hs. 307108. 0	gb:AL359553		1	3. 9	222	2. 35	0. 88	0. 37
928	Adrenal glan	217173_s_at	low density lipoprotein receptor (familial hype	Hs. 213289. 1	gb:S70123. 1		19	3. 6	268	2. 43	1. 55	0. 24
929	Adrenal glan	219655_at	hypothetical protein FLJ11808	Hs. 114611. 0	gb:NM_024728. 1	FLJ11808	7	3. 5	107	2. 03	0. 70	0. 38
930	Adrenal glan	201749_at	endothelin converting enzyme 1	Hs. 288203. 0	gb:BF969352		1	3. 4	316	2. 50	1. 81	0. 20
931	Adrenal glan	203071_at	sema domain, immunoglobulin domain (Ig), short	Hs. 82222. 0	gb:NM_004636. 1	SEMA3B	3	3. 4	364	2. 56	1. 10	0. 43
932	Adrenal glan	209237_s_at	solute carrier family 23 (nucleobase transporte	Hs. 82042. 0	gb:AF164142. 1	SVCT2	20	3. 2	223	2. 35	1. 34	0. 32
933	Adrenal glan	210469_at	discs, large (Drosophila) homolog 5	Hs. 170290. 1	gb:BC002915. 1		10	3. 2	171	2. 23	1. 64	0. 18
934	Adrenal glan	222102_at	glutathione S-transferase A3	Hs. 102484. 0	gb:NM_000847. 1	GSTA3	6	3. 2	373	2. 57	0. 94	0. 51
935	Adrenal glan	220348_at	hypothetical protein FLJ14106	Hs. 287872. 0	gb:NM_025067. 1	FLJ14106	2	2. 8	62	1. 79	0. 76	0. 37
936	Pancreas	206598_at	insulin	Hs. 89832. 0	gb:NM_000207. 1	INS	NA	17. 6	4239	3. 63	0. 29	0. 19
937	Pancreas	214377_s_at	chymotrypsin-like	Hs. 150601. 2	gb:BF508685_RC		16	17. 6	3854	3. 59	1. 80	0. 10
938	Pancreas	207062_at	islet amyloid polypeptide	Hs. 142255. 0	gb:NM_000415. 1	IAPP	12	16. 5	319	2. 50	0. 52	0. 12
939	Pancreas	207636_at	serine (or cysteine) proteinase inhibitor, cla	Hs. 158308. 0	gb:NM_006217. 2	SERPINI2	3	14	1857	3. 27	1. 65	0. 12
940	Pancreas	206297_at	chymotrypsin C (caldecrin)	Hs. 8709. 0	gb:NM_007272. 1	CTRC	1	13. 2	10148	4. 01	1. 75	0. 17
941	Pancreas	207412_x_at	carboxyl ester lipase-like (bile salt-stimulate	Hs. 169271. 0	gb:NM_001808. 1	CELL	9	13	8817	3. 95	1. 18	0. 21
942	Pancreas	220275_at	estrogen regulated gene 1	Hs. 114648. 0	gb:NM_022034. 1	ERG-1	10	10. 8	6955	3. 84	1. 65	0. 20
943	Pancreas	205910_s_at	carboxyl ester lipase (bile salt-stimulated lip	Hs. 99918. 0	gb:NM_001807. 1	CEL	9	10. 6	9467	3. 98	0. 59	0. 32
944	Pancreas	206311_s_at	phospholipase A2, group IB (pancreas)	Hs. 992. 0	gb:NM_000928. 1	PLA2G1B	12	10. 3	8738	3. 94	1. 41	0. 25
945	Pancreas	210722_at	Homo sapiens, clone MGC:12257, mRNA, complete c	Hs. 326711. 0	gb:BC005233. 1		10	9	1075	3. 03	0. 90	0. 24
946	Pancreas	207077_at	pancreatic elastase IIB	Hs. 169234. 0	gb:NM_015849. 1	LOC51032	1	8. 4	4469	3. 65	1. 02	0. 31
947	Pancreas	207778_at	rat regenerating islet-derived-like, human homo	Hs. 3191. 0	gb:NM_006508. 1	REGL	2	8. 4	3823	3. 58	1. 00	0. 31
948	Pancreas	214324_at	ESTs, Weakly similar to 2004399A chromosomal pr	Hs. 274454. 0	gb:BF222483_RC			8. 3	5096	3. 71	1. 10	0. 31
949	Pancreas	206694_at	pancreatic lipase-related protein 1	Hs. 73923. 0	gb:NM_006229. 1	PNLIPRP1	10	8. 2	9125	3. 96	1. 32	0. 32
950	Pancreas	206691_s_at	protein disulfide isomerase	Hs. 66581. 0	gb:NM_006849. 1	PDI	16	7. 8	984	2. 99	1. 31	0. 22

Fig. 66

	A	B	C	D	E	F	G	H	I	J	K	L
951	Pancreas	205615_at	carboxypeptidase A1 (pancreatic)	Hs. 2879. 0	gb:NM_001868. 1	CPA1	7	7. 6	8947	3. 95	0. 63	0. 43
952	Pancreas	206447_at	elastase 1, pancreatic	Hs. 21. 0	gb:NM_001971. 1	ELA1	1	6. 9	7171	3. 86	0. 62	0. 47
953	Pancreas	208337_s_at	nuclear receptor subfamily 5, group A, member 2	Hs. 183123. 0	gb:NM_003822. 1	NR5A2	1	5. 3	298	2. 47	1. 55	0. 17
954	Pancreas	214309_s_at	chromosome 21 open reading frame 2	Hs. 153452. 3	gb:AI435747_RC			5. 1	211	2. 32	0. 88	0. 28
955	Pancreas	211766_s_at			gb:BC005989. 1		10	5	5183	3. 71	0. 70	0. 60
956	Pancreas	221377_s_at	recombining binding protein suppressor of hair	Hs. 248217. 0	gb:NM_014276. 1	RBPSUHL	20	5	372	2. 57	1. 21	0. 27
957	Pancreas	202062_s_at	sel-1 (suppressor of lin-12, C. elegans)-like	Hs. 181300. 0	gb:NM_005065. 1	SEL1L	14	4. 3	327	2. 51	1. 71	0. 19
958	Pancreas	212942_s_at	KIAA1199 protein	Hs. 50081. 1	gb:AB033025. 1	KIAA1199	15	4. 2	353	2. 55	1. 60	0. 23
959	Pancreas	210670_at	pancreatic polypeptide	Hs. 184604. 0	gb:M15788. 1	PPY	17	4	140	2. 15	1. 46	0. 17
960	Pancreas	214446_at	ELL-RELATED RNA POLYMERASE II, ELONGATION FACT	Hs. 173334. 2	gb:NM_012081. 1	ELL2	5	3. 8	319	2. 50	1. 19	0. 35
961	Pancreas	215702_s_at	cystic fibrosis transmembrane conductance regu	Hs. 663. 1	gb:W60595_RC		7	3. 7	229	2. 36	1. 07	0. 35
962	Pancreas	202064_s_at	sel-1 (suppressor of lin-12, C. elegans)-like	Hs. 181300. 0	gb:AF052059. 1	SEL1L	14	3. 5	243	2. 38	1. 54	0. 24
963	Pancreas	219019_at	leucine rich repeat and death domain containing	Hs. 123136. 0	gb:NM_018494. 1	LRDD		3. 3	308	2. 49	1. 35	0. 34
964	Pancreas	208343_s_at	nuclear receptor subfamily 5, group A, member 2	Hs. 183123. 0	gb:AF146343. 1	CPF	1	3. 1	179	2. 25	0. 52	0. 55
965	Stomach	214476_at	trefoil factor 2 (spasmolytic protein 1)	Hs. 2979. 0	gb:NM_005423. 1	TFF2	21	23	6816	3. 83	1. 04	0. 12
966	Stomach	213265_at	Homo sapiens mRNA for caldesmon, 3 UTR	Hs. 182183. 1	gb:AI570199_RC		11	15. 6	10231	4. 01	1. 79	0. 14
967	Stomach	221095_s_at	potassium voltage-gated channel, Isk-related fa	Hs. 268538. 0	gb:NM_005136. 1	KCNE2	21	14. 7	2027	3. 31	1. 01	0. 16
968	Stomach	207546_at	ATPase, H+K+ exchanging, beta polypeptide	Hs. 813. 0	gb:NM_000705. 1	ATP4B	13	14. 5	5854	3. 77	1. 33	0. 17
969	Stomach	206334_at	lipase, gastric	Hs. 159177. 0	gb:NM_004190. 1	LIPF	10	8. 7	8467	3. 93	1. 03	0. 33
970	Stomach	207033_at	gastric intrinsic factor (vitamin B synthesis)	Hs. 110014. 0	gb:NM_005142. 1	GIF	11	8. 5	6011	3. 78	1. 15	0. 31
971	Stomach	221133_s_at	Claudin-18	Hs. 278966. 0	gb:NM_016369. 1	CLDN18	3	6. 8	384	2. 58	0. 73	0. 27
972	Stomach	207139_at	ATPase, H+K+ exchanging, alpha polypeptide	Hs. 36992. 0	gb:NM_000704. 1	ATP4A	19	6. 7	4366	3. 64	1. 37	0. 34
973	Stomach	205343_at	sulfotransferase family, cytosolic, 1C, member	Hs. 38084. 0	gb:NM_001056. 1	SULT1C1	2	6. 2	819	2. 91	1. 45	0. 24
974	Stomach	220191_at	CA11	Hs. 69319. 0	gb:NM_019617. 1	LOC56287		6. 2	9220	3. 96	0. 75	0. 52
975	Stomach	222245_s_at	hypothetical protein FLJ13459	Hs. 72222. 1	gb:AF218012. 1		20	6. 2	376	2. 58	1. 74	0. 14
976	Stomach	205199_at	carbonic anhydrase IX	Hs. 63287. 0	gb:NM_001216. 1	CA9	9	5. 4	760	2. 88	1. 67	0. 22
977	Stomach	220630_s_at	eosinophil chemotactic cytokine	Hs. 128814. 0	gb:NM_021797. 1	TSA1902	1	5. 4	1504	3. 18	1. 33	0. 34
978	Stomach	205997_at	a disintegrin and metalloproteinase domain 28	Hs. 174030. 0	gb:NM_021778. 1	ADAM28	8	4. 6	605	2. 78	0. 74	0. 44
979	Stomach	208063_s_at	calpain 9 (nCL-4)	Hs. 113292. 0	gb:NM_006615. 1	CAPN9		4. 6	249	2. 40	1. 05	0. 29
980	Stomach	214133_at	mucin 6, gastric	Hs. 159593. 0	gb:AI611214_RC			4. 5	351	2. 54	0. 79	0. 39
981	Stomach	211173_at	cholecystokinin A receptor	Hs. 129. 0	gb:D85606		4	4. 2	146	2. 16	1. 05	0. 27
982	Stomach	208269_s_at	a disintegrin and metalloproteinase domain 28	Hs. 174030. 1	gb:NM_021777. 1	ADAM28		3. 4	163	2. 21	1. 20	0. 30
983	Stomach	201467_s_at	diaphorase (NADHNADPH) (cytochrome b-5 reductas	Hs. 80706. 0	gb:AI039874_RC		16	3. 2	247	2. 39	0. 96	0. 44
984	Stomach	203884_s_at	KIAA0941 protein	Hs. 173656. 0	gb:NM_014904. 1	KIAA0941	10	3. 1	322	2. 51	1. 85	0. 21
985	Stomach	220613_s_at	hypothetical protein FLJ20163	Hs. 92254. 0	gb:NM_017695. 1	FLJ20163	11	3. 1	148	2. 17	1. 50	0. 21
986	Small Intest	207814_at	defensin, alpha 6, Paneth cell-specific	Hs. 711. 0	gb:NM_001926. 2	DEFA6	8	18. 4	4748	3. 68	0. 91	0. 15
987	Small Intest	207529_at	defensin, alpha 5, Paneth cell-specific	Hs. 72887. 0	gb:NM_021010. 1	DEFA5	8	17. 2	5310	3. 73	1. 79	0. 11
988	Small Intest	206945_at	lactase	Hs. 2251. 0	gb:NM_002299. 1	LCT	2	13. 2	2068	3. 32	1. 84	0. 11

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	A	B	C	D	E	F	G	H	I	J	K	L
989	Small Intest	210165_at	deoxyribonuclease I	Hs. 113221. 0	gb:M55983. 1	DNase I	16	10. 8	280	2. 45	0. 46	0. 18
990	Small Intest	207899_at	gastric inhibitory polypeptide	Hs. 1454. 0	gb:NM_004123. 1	GIP		9. 5	1032	3. 01	0. 85	0. 23
991	Small Intest	211618_s_at			gb:M31008. 1	ALPI	2	8. 8	101	2. 00	0. 24	0. 20
992	Small Intest	207259_at	hypothetical protein FLJ20694	Hs. 232802. 0	gb:NM_017928. 1	FLJ20694	17	8. 2	81	1. 91	0. 20	0. 21
993	Small Intest	207885_at	calbindin 3, (vitamin D-dependent calcium-bindi	Hs. 639. 0	gb:NM_004057. 1	CALB3	X	8. 2	200	2. 30	0. 72	0. 19
994	Small Intest	220639_at	hypothetical protein FLJ22800	Hs. 156652. 0	gb:NM_024795. 1	FLJ22800		7. 4	806	2. 91	0. 92	0. 27
995	Small Intest	206894_at	apolipoprotein A-IV	Hs. 1247. 0	gb:NM_000482. 2	APOA4	11	7. 1	1360	3. 13	0. 49	0. 37
996	Small Intest	207158_at	apolipoprotein B mRNA editing enzyme, catalytic	Hs. 560. 0	gb:NM_001644. 2	APOBEC1	12	7. 1	332	2. 52	1. 60	0. 13
997	Small Intest	219640_at	claudin 15	Hs. 38738. 0	gb:NM_014343. 1	CLDN15	7	6. 1	1116	3. 05	1. 55	0. 25
998	Small Intest	207378_at	trehalase (brush-border membrane glycoprotein)	Hs. 129712. 0	gb:NM_007180. 1	TREH	11	5. 5	133	2. 12	0. 28	0. 34
999	Small Intest	207251_at	meprin A, beta	Hs. 194777. 0	gb:NM_005925. 1	MEP1B	18	5. 1	1043	3. 02	1. 41	0. 31
1000	Small Intest	207638_at	protease, serine, 7 (enterokinase)	Hs. 158333. 0	gb:NM_002772. 1	PRSS7	21	5	393	2. 59	0. 78	0. 36
1001	Small Intest	220074_at	mucin and cadherin-like	Hs. 165619. 1	gb:NM_017717. 2	MUCDHL	NA	4. 9	454	2. 66	0. 79	0. 38
1002	Small Intest	206988_at	small inducible cytokine subfamily A (Cys-Cys),	Hs. 50404. 0	gb:NM_005624. 1	SCYA25	19	4. 8	793	2. 90	1. 25	0. 34
1003	Small Intest	207473_at	motilin	Hs. 2813. 0	gb:NM_002418. 1	MLN	6	4. 6	250	2. 40	1. 71	0. 15
1004	Small Intest	211427_s_at	potassium inwardly-rectifying channel, subfamily	Hs. 296361. 0	gb:AB013889. 1			4. 6	76	1. 88	0. 39	0. 33
1005	Small Intest	220186_s_at	hypothetical protein FLJ20124	Hs. 4205. 0	gb:NM_017675. 1	FLJ20124	5	4. 5	424	2. 63	1. 73	0. 20
1006	Small Intest	220106_at	NPC1 (Niemann-Pick disease, type C1, gene)-like	Hs. 47701. 0	gb:NM_013389. 1	NPC1L1	7	4. 4	352	2. 55	1. 19	0. 31
1007	Small Intest	220421_at	hypothetical protein FLJ21458	Hs. 189109. 0	gb:NM_024850. 1	FLJ21458	5	4. 4	252	2. 40	0. 92	0. 34
1008	Small Intest	219614_s_at	X transporter protein 3	Hs. 107854. 0	gb:NM_020208. 1	XT3	3	4. 2	155	2. 19	0. 86	0. 32
1009	Small Intest	208323_s_at	annexin A13	Hs. 181107. 0	gb:NM_004306. 1	ANXA13	8	4. 1	346	2. 54	1. 30	0. 31
1010	Small Intest	209749_s_at	angiotensin I converting enzyme (peptidyl-dipep	Hs. 298469. 1	gb:AI623989_RC		17	4	131	2. 12	0. 86	0. 31
1011	Small Intest	211184_s_at	PDZ-73 protein	Hs. 132945. 2	gb:AB006955. 1	aie-75	11	4	310	2. 49	1. 59	0. 22
1012	Small Intest	210177_at	ring finger protein 9	Hs. 274295. 0	gb:U34249. 1	ZNFB7	6	3. 9	103	2. 01	0. 82	0. 31
1013	Small Intest	207254_at	solute carrier family 15 (oligopeptide transpor	Hs. 2217. 0	gb:NM_005073. 1	SLC15A1	13	3. 6	233	2. 37	0. 58	0. 50
1014	Small Intest	207567_at	solute carrier family 13 (sodium-dependent dica	Hs. 102307. 0	gb:NM_003984. 1	SLC13A2	17	3. 6	109	2. 04	0. 76	0. 35
1015	Small Intest	206484_s_at	X-prolyl aminopeptidase (aminopeptidase P) 2, m	Hs. 57922. 0	gb:NM_003399. 3	XPNPEP2	X	3. 5	425	2. 63	1. 43	0. 34
1016	Small Intest	206964_at	putative N-acetyltransferase Camello 2	Hs. 278983. 0	gb:NM_016347. 1	CML2	2	3. 4	155	2. 19	1. 31	0. 26
1017	Small Intest	217054_at	mucin 3A, intestinal	Hs. 129782. 2	gb:AF007194. 1_RC	MUC3		3. 3	67	1. 83	0. 93	0. 27
1018	Small Intest	211303_x_at	folate hydrolase (prostate-specific membrane ar	Hs. 1915. 1	gb:AF261715. 1	PSMALGCP I	11	3. 2	628	2. 80	1. 02	0. 56
1019	Small Intest	219332_at	hypothetical protein FLJ23471	Hs. 110964. 0	gb:NM_024723. 1	FLJ23471		3. 1	276	2. 44	1. 72	0. 23
1020	Small Intest	207456_at	hepatocyte nuclear factor 4, gamma	Hs. 241529. 0	gb:NM_004133. 2	HNF4G	8	3	72	1. 85	0. 83	0. 34
1021	Small Intest	210179_at	potassium inwardly-rectifying channel, subfamily	Hs. 296361. 1	gb:AJ007557. 1	Kir7. 1	2	2. 6	84	1. 92	0. 85	0. 41
1022	Colon	211848_s_at	carcinoembryonic antigen-related cell adhesion	Hs. 74466. 1	gb:AF006623. 1	CGM2	19	9. 5	832	2. 92	0. 72	0. 23
1023	Colon	220645_at	hypothetical protein FLJ20127	Hs. 179100. 0	gb:NM_017678. 1	FLJ20127	11	8. 8	452	2. 66	0. 88	0. 20
1024	Colon	220834_at	hypothetical protein FLJ20217	Hs. 272789. 0	gb:NM_017716. 1	FLJ20217	11	7. 6	970	2. 99	1. 50	0. 20
1025	Colon	206268_at	left-right determination, factor B	Hs. 278239. 0	gb:NM_020997. 1	LEFTB	1	7. 1	750	2. 87	1. 72	0. 16
1026	Colon	206418_at	NADPH oxidase 1	Hs. 132370. 1	gb:NM_007052. 2	NOX1	X	6. 8	157	2. 20	0. 27	0. 28

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	A	B	C	D	E	F	G	H	I	J	K	L
1027	Colon	205593_s_at	phosphodiesterase 9A	Hs. 18953. 0	gb:NM_002606. 1	PDE9A	21	6. 4	280	2. 45	0. 84	0. 25
1028	Colon	220026_at	chloride channel, calcium activated, family mer	Hs. 227059. 0	gb:NM_012128. 2	CLCA4	1	6. 3	1515	3. 18	1. 27	0. 30
1029	Colon	206198_s_at	carcinoembryonic antigen-related cell adhesion	Hs. 74466. 0	gb:L31792. 1	CGM2	19	5. 8	786	2. 90	1. 01	0. 33
1030	Colon	215657_at	Homo sapiens cDNA: FLJ21391 fis, clone COL0347	Hs. 306752. 0	gb:AK025044. 1_RC		7	4. 9	304	2. 48	0. 50	0. 40
1031	Colon	207432_at	hypothetical protein FLJ20132	Hs. 190222. 0	gb:NM_017682. 1	FLJ20132	19	4. 7	642	2. 81	1. 18	0. 34
1032	Colon	207080_s_at	peptide YY	Hs. 169249. 0	gb:NM_004160. 1	PYY	17	3. 6	458	2. 66	1. 05	0. 44
1033	Colon	206208_at	carbonic anhydrase IV	Hs. 89485. 0	gb:NM_000717. 2	CA4	17	3. 2	331	2. 52	1. 18	0. 41
1034	Colon	221091_at	insulin-like 5	Hs. 251380. 0	gb:NM_005478. 2	INSL5	1	2. 6	125	2. 10	1. 21	0. 34
1035	Bladder	205127_at	prostaglandin-endoperoxide synthase 1 (prostag	Hs. 88474. 0	gb:NM_000962. 1	PTGS1	9	5. 5	551	2. 74	1. 19	0. 28
1036	Bladder	33760_at						4. 9	164	2. 21	1. 75	0. 10
1037	Bladder	214624_at	uroplakin 1A	Hs. 159309. 0	gb:AA548647_RC		19	4. 6	612	2. 79	1. 49	0. 28
1038	Bladder	209298_s_at	intersectin 1 (SH3 domain protein)	Hs. 66392. 1	gb:AF114488. 1	ITSN	21	4. 5	182	2. 26	1. 46	0. 18
1039	Bladder	206463_s_at	short-chain alcohol dehydrogenase family membe	Hs. 272499. 0	gb:NM_005794. 1	HEP27	14	4. 1	904	2. 96	1. 17	0. 43
1040	Bladder	211819_s_at	SH3-domain protein 5 (ponsin)	Hs. 108924. 2	gb:AF136381. 1	SH3P12		4. 1	267	2. 43	1. 67	0. 18
1041	Bladder	216030_s_at	semenogelin II	Hs. 180016. 1	gb:AL049767		20	4. 1	272	2. 43	0. 76	0. 41
1042	Bladder	212806_at	KIAA0367 protein	Hs. 23311. 0	gb:AL138349_RC		9	3. 8	171	2. 23	1. 36	0. 23
1043	Bladder	211562_s_at	leiomodin 1 (smooth muscle)	Hs. 79386. 1	gb:BC001755. 1			3. 3	214	2. 33	1. 00	0. 41
1044	Bladder	203813_s_at	slit (Drosophila) homolog 3	Hs. 57929. 0	gb:NM_003062. 1	SLIT3	5	2. 4	216	2. 34	1. 30	0. 43
1045	Prostate	209854_s_at	kallikrein 2, prostatic	Hs. 181350. 0	gb:AA595465_RC		19	11. 3	836	2. 92	0. 74	0. 19
1046	Prostate	204582_s_at	kallikrein 3, (prostate specific antigen)	Hs. 171995. 0	gb:NM_001648. 1	KLK3	19	7. 6	2608	3. 42	0. 74	0. 35
1047	Prostate	204393_s_at	acid phosphatase, prostate	Hs. 1852. 0	gb:NM_001099. 2	ACPP	3	6. 5	2118	3. 33	1. 60	0. 26
1048	Prostate	204583_x_at	kallikrein 3, (prostate specific antigen)	Hs. 171995. 0	gb:U17040. 1		19	6. 1	2775	3. 44	1. 08	0. 39
1049	Prostate	208180_s_at	H4 histone family, member H	Hs. 93758. 0	gb:NM_003543. 2	H4FH	6	4. 6	161	2. 21	1. 41	0. 17
1050	Prostate	210339_s_at	kallikrein 2, prostatic	Hs. 181350. 1	gb:BC005196. 1		19	4. 5	565	2. 75	1. 52	0. 27
1051	Prostate	206569_at	suppression of tumorigenicity 16 (melanoma dif	Hs. 315463. 0	gb:NM_006850. 1	ST16	1	4. 1	92	1. 96	0. 69	0. 31
1052	Prostate	205940_at	myosin, heavy polypeptide 3, skeletal muscle,	Hs. 173084. 0	gb:NM_002470. 1	MYH3	17	3. 7	195	2. 29	0. 59	0. 46
1053	Testis	215622_x_at	hypothetical protein	Hs. 30127. 3	gb:AL137671. 1	DKFZp434L1	3	17. 8	2601	3. 42	1. 93	0. 08
1054	Testis	210122_at	protamine 2	Hs. 2324. 0	gb:BC005303. 1		16	16. 6	3636	3. 56	0. 32	0. 20
1055	Testis	207344_at	A kinase (PRKA) anchor protein 3	Hs. 98397. 0	gb:NM_006422. 1	AKAP3	12	15. 3	518	2. 71	0. 95	0. 12
1056	Testis	210415_s_at	outer dense fibre of sperm tails 2	Hs. 129055. 1	gb:AF053970. 1		9	14. 2	1297	3. 11	0. 50	0. 18
1057	Testis	214183_s_at	transketolase-like 1	Hs. 102866. 1	gb:X91817. 1		X	13. 8	984	2. 99	0. 76	0. 16
1058	Testis	221630_s_at	VASA protein	Hs. 97689. 1	gb:AY004154. 1	VASA	5	13. 7	787	2. 90	0. 20	0. 20
1059	Testis	206406_at	mitochondrial capsule selenoprotein	Hs. 111850. 0	gb:NM_030663. 1	MCSP	1	13. 3	1254	3. 10	0. 87	0. 17
1060	Testis	220299_at	hypothetical protein	Hs. 143251. 0	gb:NM_019073. 1	FLJ10007	1	12. 7	995	3. 00	1. 63	0. 11
1061	Testis	220501_at	actin-like 7A	Hs. 123530. 0	gb:NM_006687. 2	ACTL7A	9	12. 4	401	2. 60	0. 81	0. 14
1062	Testis	206587_at	chaperonin containing TCP1, subunit 6B (zeta 2)	Hs. 73072. 0	gb:NM_006584. 1	CCT6B	17	12. 2	940	2. 97	1. 77	0. 10
1063	Testis	206319_s_at	eppin-3	Hs. 121084. 0	gb:AF286368. 1	EPPIN1	20	11. 6	709	2. 85	1. 57	0. 11
1064	Testis	207336_at	SRY (sex determining region Y)-box 5	Hs. 87224. 0	gb:NM_006940. 1	SOX5	8	11. 6	155	2. 19	0. 53	0. 14

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	A	B	C	D	E	F	G	H	I	J	K	L
1065	Testis	207918_s_at	testis specific protein, Y-linked	Hs. 2051. 0	gb:NM_003308. 2	TSPY	Y	11. 5	493	2. 69	0. 21	0. 22
1066	Testis	220922_s_at	sperm protein associated with the nucleus, X cl	Hs. 293266. 0	gb:NM_013453. 1	SPANXA1	X	11. 4	507	2. 70	0. 82	0. 17
1067	Testis	208235_x_at	G antigen 7	Hs. 278606. 0	gb:NM_021123. 1	GAGE7		10. 4	929	2. 97	1. 32	0. 16
1068	Testis	207780_at	cylicin, basic protein of sperm head cytoskelet	Hs. 3232. 0	gb:NM_001340. 1	CYLC2	9	10. 3	392	2. 59	1. 45	0. 11
1069	Testis	206358_at	protamine 1	Hs. 2909. 0	gb:NM_002761. 1	PRM1	16	10. 1	4920	3. 69	0. 87	0. 28
1070	Testis	203015_s_at	KIAA0923 protein	Hs. 22587. 0	gb:AW136988_RC		1	10	570	2. 76	1. 83	0. 09
1071	Testis	219820_at	NTT5 protein	Hs. 59260. 0	gb:NM_014037. 1	NTT5	19	10	808	2. 91	1. 71	0. 12
1072	Testis	219928_s_at	fibrousheathin II	Hs. 314452. 0	gb:NM_012189. 1	FSP-2	18	9. 8	1080	3. 03	1. 77	0. 13
1073	Testis	220378_at	t-complex 11 (a murine tcp homolog)	Hs. 128322. 0	gb:NM_018679. 1	TCP11	6	9. 7	2926	3. 47	1. 61	0. 19
1074	Testis	210497_x_at	synovial sarcoma, X breakpoint 2	Hs. 289105. 2	gb:BC002818. 1		X	9. 6	401	2. 60	1. 13	0. 15
1075	Testis	219484_at	host cell factor 2	Hs. 55601. 0	gb:NM_013320. 1	HCF-2	12	9. 6	398	2. 60	1. 63	0. 10
1076	Testis	220057_at	XAGE-1 protein	Hs. 112208. 0	gb:NM_020411. 1	XAGE-1	X	9. 6	53	1. 72	0. 28	0. 15
1077	Testis	207973_x_at	acrosomal vesicle protein 1	Hs. 169222. 10	gb:NM_020110. 1	ACRV1	11	9. 4	929	2. 97	1. 91	0. 11
1078	Testis	215430_at	H. sapiens mRNA for glycerol kinase testis speci	Hs. 98008. 0	gb:AA757089_RC		4	9. 4	1058	3. 02	1. 59	0. 15
1079	Testis	220271_x_at	hypothetical protein FLJ23588	Hs. 99330. 0	gb:NM_022785. 1	FLJ23588	22	9. 4	108	2. 03	0. 63	0. 15
1080	Testis	221611_s_at	hypothetical protein	Hs. 30127. 0	gb:AY014283. 1		3	9. 4	1705	3. 23	1. 28	0. 21
1081	Testis	207377_at	type 1 protein phosphatase inhibitor	Hs. 127689. 0	gb:NM_025210. 1	I-4	X	9. 3	150	2. 18	0. 44	0. 19
1082	Testis	214485_at	outer dense fibre of sperm tails 1	Hs. 159274. 0	gb:NM_024410. 1	ODF1	8	9. 2	1424	3. 15	1. 56	0. 17
1083	Testis	219587_at	hypothetical protein FLJ20535	Hs. 183362. 0	gb:NM_017868. 1	FLJ20535	11	9. 2	220	2. 34	0. 55	0. 19
1084	Testis	206776_x_at	acrosomal vesicle protein 1	Hs. 169222. 4	gb:NM_001612. 2	ACRV1	11	9. 1	819	2. 91	1. 92	0. 11
1085	Testis	211107_s_at	serinethreonine kinase 13 (auroraIPL1-like)	Hs. 98338. 1	gb:AB017332. 1	aik3	19	9	390	2. 59	1. 84	0. 08
1086	Testis	213124_at	DKFZP434N043 protein	Hs. 59255. 0	gb:BG538800		19	9	438	2. 64	1. 72	0. 10
1087	Testis	215320_at	DKFZP434M131 protein	Hs. 189296. 0	gb:AL080179. 1	DKFZp434M1	9	9	38	1. 58	-0. 06	0. 18
1088	Testis	208155_x_at	G antigen 6	Hs. 272484. 0	gb:NM_001476. 1	GAGE6	X	8. 8	1045	3. 02	1. 53	0. 17
1089	Testis	220205_at	transmembrane phosphatase with tensin homology	Hs. 122986. 0	gb:NM_013315. 1	TPTE	21	8. 7	1373	3. 14	1. 02	0. 24
1090	Testis	205406_s_at	sperm autoantigenic protein 17	Hs. 286233. 0	gb:NM_017425. 1	SPA17	11	8. 6	854	2. 93	1. 61	0. 15
1091	Testis	216663_s_at	BLu protein	Hs. 167380. 2	gb:AC002481			8. 6	612	2. 79	1. 73	0. 12
1092	Testis	203194_s_at	nucleoporin 98kD	Hs. 112255. 0	gb:AA527238_RC		11	8. 5	562	2. 75	1. 83	0. 11
1093	Testis	207086_x_at	G antigen 4	Hs. 183199. 0	gb:NM_001474. 1	GAGE4	X	8. 5	1078	3. 03	1. 58	0. 17
1094	Testis	211659_at			gb:M76676. 1		14	8. 5	225	2. 35	0. 53	0. 21
1095	Testis	205702_at	putative homeodomain transcription factor	Hs. 123637. 0	gb:NM_006608. 1	PHTF1	1	8. 4	328	2. 52	1. 87	0. 08
1096	Testis	206740_x_at	synaptonemal complex protein 1	Hs. 112743. 0	gb:NM_003176. 1	SYCP1	1	8. 4	407	2. 61	1. 50	0. 13
1097	Testis	220118_at	testis zinc finger protein	Hs. 99430. 0	gb:NM_014383. 1	TZFP	19	8. 4	178	2. 25	0. 66	0. 19
1098	Testis	220179_at	putative metallopeptidase (family M19)	Hs. 302028. 0	gb:NM_022357. 1	LOC64180	16	8. 3	267	2. 43	0. 65	0. 21
1099	Testis	222106_at	prion gene complex, downstream	Hs. 121281. 1	gb:AL133396		20	8. 3	1077	3. 03	1. 43	0. 19
1100	Testis	207990_x_at	acrosomal vesicle protein 1	Hs. 169222. 5	gb:NM_020114. 1	ACRV1	11	8. 2	648	2. 81	1. 95	0. 11
1101	Testis	213370_s_at	DKFZP434L243 protein	Hs. 21695. 0	gb:BF057298_RC		3	8. 2	494	2. 69	1. 85	0. 10
1102	Testis	220422_at	ubiquilin 3	Hs. 189184. 0	gb:NM_017481. 1	UBQLN3	11	8. 2	778	2. 89	1. 50	0. 17

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	A	B	C	D	E	F	G	H	I	J	K	L
1103	Testis	210712_at	Homo sapiens lactate dehydrogenase A mRNA, comp	Hs. 307052. 0	gb:AY009108. 1		15	8. 1	539	2. 73	1. 35	0. 17
1104	Testis	213454_at	cortistatin	Hs. 44205. 1	gb:AL578583_RC		1	8. 1	268	2. 43	1. 72	0. 09
1105	Testis	207991_x_at	acrosomal vesicle protein 1	Hs. 169222. 6	gb:NM_020113. 1	ACRV1	11	8	561	2. 75	1. 91	0. 11
1106	Testis	217009_at	phosphoglycerate kinase 1	Hs. 78771. 2	gb:AL121974			8	594	2. 77	1. 32	0. 18
1107	Testis	220110_s_at	nuclear RNA export factor 3	Hs. 60386. 0	gb:NM_022052. 1	NXF3	X	8	246	2. 39	0. 41	0. 25
1108	Testis	220498_at	actin-like 7B	Hs. 119287. 0	gb:NM_006686. 2	ACTL7B	9	7. 9	730	2. 86	1. 06	0. 23
1109	Testis	207736_s_at	transition protein 2 (during histone to protami	Hs. 2748. 0	gb:NM_005425. 1	TNP2	16	7. 8	70	1. 85	0. 31	0. 20
1110	Testis	205408_at	myeloidlymphoid or mixed-lineage leukemia (tri	Hs. 301209. 0	gb:NM_004641. 1	AF10	10	7. 7	426	2. 63	1. 90	0. 09
1111	Testis	221035_s_at			gb:NM_031272. 1	TEX14		7. 6	381	2. 58	0. 89	0. 22
1112	Testis	207022_s_at	lactate dehydrogenase C	Hs. 99881. 0	gb:NM_002301. 2	LDHC	11	7. 5	1564	3. 19	0. 90	0. 31
1113	Testis	207232_s_at	KIAA0675 gene product	Hs. 165662. 0	gb:NM_014648. 1	KIAA0675		7. 5	123	2. 09	0. 74	0. 18
1114	Testis	206640_x_at	G antigen 7B	Hs. 251677. 0	gb:NM_001477. 1	GAGE7B	X	7. 4	961	2. 98	1. 25	0. 23
1115	Testis	211403_x_at	variable charge, X chromosome	Hs. 278906. 1	gb:AF167079. 1	VCXB	X	7. 4	311	2. 49	1. 44	0. 14
1116	Testis	206733_at	tubby like protein 2	Hs. 104636. 0	gb:NM_003323. 1	TULP2	19	7. 3	281	2. 45	1. 70	0. 10
1117	Testis	210191_s_at	putative homeodomain transcription factor	Hs. 123637. 1	gb:BC002447. 1		1	7. 3	342	2. 53	1. 81	0. 10
1118	Testis	220379_at	fascin (Strongylocentrotus purpuratus) homolog	Hs. 128402. 0	gb:NM_020369. 1	FSCN3	7	7. 3	801	2. 90	1. 90	0. 14
1119	Testis	215734_at	Homo sapiens chromosome 19, cosmid R26660	Hs. 90010. 1	gb:AW182303_RC		NA	7. 2	1490	3. 17	0. 87	0. 32
1120	Testis	220256_s_at	hypothetical protein FKSG25	Hs. 302023. 0	gb:NM_022120. 1	FLJ00030	1	7. 2	207	2. 32	0. 97	0. 19
1121	Testis	214397_at	methyl-CpG binding domain protein 2	Hs. 25674. 2	gb:AI827820_RC		18	7. 1	272	2. 43	1. 34	0. 15
1122	Testis	208013_s_at	acrosomal vesicle protein 1	Hs. 169222. 9	gb:NM_020115. 1	ACRV1	11	7	527	2. 72	1. 81	0. 13
1123	Testis	214264_s_at	Homo sapiens, clone MGC:2492, mRNA, complete cd	Hs. 87819. 1	gb:AI656610_RC		14	7	479	2. 68	1. 64	0. 15
1124	Testis	221337_s_at	a disintegrin and metalloproteinase domain 29	Hs. 126838. 1	gb:NM_021780. 1	ADAM29		7	268	2. 43	1. 08	0. 19
1125	Testis	214409_at	ret finger protein-like 3 antisense	Hs. 274285. 0	gb:AL021937		22	6. 9	350	2. 54	1. 36	0. 17
1126	Testis	207797_s_at	hypothetical protein DKFZp76100113	Hs. 42768. 0	gb:NM_018409. 1	DKFZp76100	4	6. 8	177	2. 25	0. 79	0. 21
1127	Testis	219128_at	hypothetical protein FLJ20558	Hs. 38681. 0	gb:NM_017880. 1	FLJ20558	2	6. 8	493	2. 69	1. 88	0. 12
1128	Testis	220426_at	hypothetical protein MGC5356	Hs. 197755. 0	gb:NM_024059. 1	MGC5356	20	6. 8	590	2. 77	1. 54	0. 18
1129	Testis	213220_at	Homo sapiens cDNA: FLJ22071 fis, clone HEP11691	Hs. 326248. 2	gb:AV706096		10	6. 7	539	2. 73	1. 70	0. 15
1130	Testis	219784_at	hypothetical protein FLJ22477	Hs. 118944. 0	gb:NM_024735. 1	FLJ22477	16	6. 7	331	2. 52	1. 16	0. 20
1131	Testis	221194_s_at	PTD016 protein	Hs. 30154. 0	gb:NM_016125. 1	LOC51136	17	6. 7	367	2. 56	1. 77	0. 12
1132	Testis	212691_at	SH3-containing protein SH3GLB2	Hs. 30002. 3	gb:AW131863_RC		9	6. 6	520	2. 72	2. 03	0. 10
1133	Testis	213346_at	Homo sapiens clone 24463 mRNA sequence	Hs. 13413. 0	gb:BE748563_RC		13	6. 6	620	2. 79	1. 77	0. 15
1134	Testis	218586_at	hypothetical protein FLJ10914	Hs. 143954. 0	gb:NM_018270. 1	FLJ10914	20	6. 6	251	2. 40	1. 70	0. 11
1135	Testis	210352_at	thyroid hormone receptor coactivating protein	Hs. 5464. 1	gb:AL136823. 1	DKFZp434P2	5	6. 5	795	2. 90	1. 32	0. 24
1136	Testis	214672_at	KIAA0998 protein	Hs. 131525. 2	gb:AB023215. 1	KIAA0998	14	6. 5	453	2. 66	1. 76	0. 14
1137	Testis	218526_s_at	HSPC165 protein	Hs. 13605. 0	gb:NM_014185. 1	HSPC165	17	6. 5	391	2. 59	2. 00	0. 09
1138	Testis	206112_at	testis-specific ankyrin motif containing protei	Hs. 73073. 0	gb:NM_019644. 1	LOC56311	7	6. 4	1959	3. 29	0. 43	0. 45
1139	Testis	207116_s_at	glyceraldehyde-3-phosphate dehydrogenase, testi	Hs. 248017. 0	gb:NM_014364. 1	GAPDS	19	6. 4	982	2. 99	1. 23	0. 27
1140	Testis	220284_at	soggy-1 gene	Hs. 124021. 0	gb:NM_014419. 1	SGY-1	19	6. 4	370	2. 57	0. 73	0. 28

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	A	B	C	D	E	F	G	H	I	J	K	L
1141	Testis	207208_at	testes-specific heterogenous nuclear ribonucle	Hs. 121605. 0	gb:NM_014469. 2	HNRNPG-T		6. 3	231	2. 36	1. 16	0. 19
1142	Testis	209698_at	HCR (a-helix coiled-coil rod homologue)	Hs. 110746. 0	gb:AF216493. 1	HCR	6	6. 3	410	2. 61	1. 24	0. 22
1143	Testis	220744_s_at	WD repeat domain 10	Hs. 70202. 0	gb:NM_018262. 1	FLJ10897	3	6. 3	680	2. 83	0. 96	0. 30
1144	Testis	201823_s_at	ring finger protein 14	Hs. 215857. 0	gb:NM_004290. 1	RNF14	5	6. 2	298	2. 47	1. 94	0. 09
1145	Testis	206608_s_at	retinitis pigmentosa GTPase regulator interacti	Hs. 131917. 0	gb:NM_020366. 1	RPGRIP1	14	6. 2	305	2. 48	1. 70	0. 13
1146	Testis	206719_at	synaptogyrin 4	Hs. 120857. 0	gb:NM_012451. 1	SYNGR4	19	6. 2	280	2. 45	1. 52	0. 15
1147	Testis	222239_s_at	deleted in cancer 1; RNA helicase HDBDICE1	Hs. 58570. 1	gb:AL117626. 1	DKFZp434B1	13	6. 2	202	2. 31	1. 49	0. 13
1148	Testis	206568_at	transition protein 1 (during histone to protami	Hs. 3017. 0	gb:NM_003284. 1	TNP1	2	6. 1	3404	3. 53	0. 89	0. 43
1149	Testis	207271_x_at	sperm associated antigen 11	Hs. 2717. 0	gb:NM_016512. 1	HE2	8	6. 1	174	2. 24	0. 65	0. 26
1150	Testis	215142_at	Huntingtin interacting protein M	Hs. 122959. 0	gb:AA815276_RC		X	6. 1	221	2. 34	0. 87	0. 24
1151	Testis	209922_at	BRCA1 associated protein	Hs. 122764. 0	gb:AF035620. 1	BRAP2	12	6	152	2. 18	0. 34	0. 30
1152	Testis	218883_s_at	hypothetical protein FLJ23468	Hs. 38178. 0	gb:NM_024629. 1	FLJ23468	4	6	1040	3. 02	1. 69	0. 22
1153	Testis	220239_at	SBBI26 protein	Hs. 26481. 0	gb:NM_018846. 1	SBBI26	7	6	407	2. 61	1. 51	0. 18
1154	Testis	210642_at	calicin	Hs. 115460. 0	gb:AF333334. 1		9	5. 9	248	2. 39	1. 00	0. 24
1155	Testis	214518_at	pyruvate dehydrogenase (lipoamide) alpha 2	Hs. 131361. 0	gb:NM_005390. 1	PDHA2	4	5. 9	858	2. 93	1. 49	0. 25
1156	Testis	216471_x_at	synovial sarcoma, X breakpoint 2	Hs. 289105. 5	gb:X79200. 1		X	5. 9	193	2. 28	1. 19	0. 19
1157	Testis	220298_s_at	hypothetical protein	Hs. 143251. 0	gb:NM_019073. 1	FLJ10007	1	5. 9	804	2. 91	0. 80	0. 36
1158	Testis	220769_s_at	hypothetical protein FLJ23129	Hs. 130401. 0	gb:NM_024763. 1	FLJ23129	1	5. 9	174	2. 24	1. 54	0. 12
1159	Testis	222351_at	ESTs, Highly similar to 2AAB_HUMAN SERINETHREON	Hs. 168737. 0	gb:AW009884_RC		11	5. 9	239	2. 38	0. 56	0. 31
1160	Testis	204634_at	serinethreonine kinase 2	Hs. 1087. 0	gb:NM_003157. 1	STK2	3	5. 8	372	2. 57	1. 78	0. 14
1161	Testis	206218_at	melanoma antigen, family B, 2	Hs. 113824. 0	gb:NM_002364. 1	MAGEB2	X	5. 8	224	2. 35	1. 52	0. 14
1162	Testis	207020_at	heat shock transcription factor 2 binding prote	Hs. 97624. 0	gb:NM_007031. 1	HSF2BP	21	5. 8	246	2. 39	1. 81	0. 10
1163	Testis	209378_s_at	KIAA1128 protein	Hs. 81897. 0	gb:AI478879_RC		10	5. 8	731	2. 86	1. 73	0. 20
1164	Testis	220107_s_at	hypothetical protein FLJ23093	Hs. 48642. 0	gb:NM_024643. 1	FLJ23093	14	5. 8	384	2. 58	1. 22	0. 24
1165	Testis	1255_g_at						5. 7	169	2. 23	1. 34	0. 16
1166	Testis	203521_s_at	endocrine regulator	Hs. 48433. 0	gb:NM_014345. 1	HRIHFB2436	6	5. 7	304	2. 48	1. 73	0. 13
1167	Testis	204109_s_at	nuclear transcription factor Y, alpha	Hs. 797. 1	gb:NM_002505. 2	NFYA	6	5. 7	216	2. 33	1. 77	0. 10
1168	Testis	204190_at	highly charged protein	Hs. 151236. 0	gb:NM_005800. 1	D13S106E	13	5. 7	279	2. 45	1. 98	0. 08
1169	Testis	204556_s_at	KIAA0996 protein	Hs. 60177. 0	gb:AL568422_RC		13	5. 7	540	2. 73	1. 65	0. 19
1170	Testis	207019_s_at	A kinase (PRKA) anchor protein 4	Hs. 97633. 0	gb:NM_003886. 1	AKAP4		5. 7	506	2. 70	0. 87	0. 32
1171	Testis	211519_s_at	kinesin-like 6 (mitotic centromere-associated k	Hs. 69360. 0	gb:AY026505. 1			5. 7	405	2. 61	1. 79	0. 14
1172	Testis	212272_at	lipin 1	Hs. 81412. 0	gb:AA813260_RC		2	5. 7	511	2. 71	1. 50	0. 21
1173	Testis	214147_at	Homo sapiens mRNA; cDNA DKFZp434G153 (from clor	Hs. 225433. 0	gb:AL046350_RC		1	5. 7	320	2. 51	1. 28	0. 21
1174	Testis	214400_at	insulin-like 3 (Leydig cell)	Hs. 37062. 0	gb:AI991694_RC		19	5. 7	1432	3. 16	0. 64	0. 44
1175	Testis	213322_at	Human DNA sequence from clone 34B21 on chromosc	Hs. 183056. 0	gb:AL031778		6	5. 6	240	2. 38	1. 85	0. 10
1176	Testis	214371_at	serinethreonine kinase 22B (spermiogenesis assc	Hs. 103978. 0	gb:AI652441_RC		22	5. 6	688	2. 84	1. 49	0. 24
1177	Testis	214418_at	Homo sapiens mRNA; cDNA DKFZp434D024 (from clor	Hs. 97823. 0	gb:AI656822_RC		15	5. 6	168	2. 23	0. 48	0. 31
1178	Testis	221716_s_at	hypothetical protein PRTD-NY3	Hs. 99344. 0	gb:AY009107. 1		19	5. 6	777	2. 89	1. 66	0. 22

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	A	B	C	D	E	F	G	H	I	J	K	L
1179	Testis	203135_at	TATA box binding protein	Hs. 1100. 0	gb:NM_003194. 1	TBP	6	5. 5	251	2. 40	1. 78	0. 11
1180	Testis	208107_s_at			gb:NM_030941. 1	LOC81691	16	5. 5	2137	3. 33	1. 22	0. 38
1181	Testis	210236_at	protein tyrosine phosphatase, receptor type, f	Hs. 183648. 1	gb:U22815. 1		11	5. 5	205	2. 31	1. 29	0. 18
1182	Testis	210441_at	CDC14 (cell division cycle 14, S. cerevisiae) h	Hs. 65993. 1	gb:AF064102. 1		1	5. 5	349	2. 54	1. 18	0. 25
1183	Testis	212201_at	KIAA0692 protein	Hs. 100729. 0	gb:AW274877_RC			5. 5	374	2. 57	1. 72	0. 16
1184	Testis	220973_s_at			gb:NM_030974. 1	DKFZP434N1923		5. 5	105	2. 02	0. 97	0. 19
1185	Testis	220996_s_at			gb:NM_030933. 1	GE36		5. 5	576	2. 76	1. 07	0. 31
1186	Testis	205543_at	heat shock protein (hsp110 family)	Hs. 71992. 0	gb:NM_014278. 1	APG-1	4	5. 4	777	2. 89	1. 45	0. 27
1187	Testis	206763_at	FK506-binding protein 6 (36kD)	Hs. 150490. 0	gb:NM_003602. 1	FKBP6	7	5. 4	235	2. 37	1. 46	0. 17
1188	Testis	215215_s_at	Homo sapiens Chromosome 16 BAC clone CIT987SK-4	Hs. 307353. 0	gb:AC004381		16	5. 4	1865	3. 27	1. 03	0. 42
1189	Testis	218349_s_at	hypothetical protein FLJ10036	Hs. 21331. 0	gb:NM_017975. 1	FLJ10036	15	5. 4	277	2. 44	0. 69	0. 33
1190	Testis	221037_s_at			gb:NM_031291. 1	DKFZP434N1235		5. 4	359	2. 56	0. 64	0. 36
1191	Testis	222064_s_at	hypothetical protein MGC2744	Hs. 317403. 1	gb:AI093187_RC		17	5. 4	224	2. 35	1. 75	0. 11
1192	Testis	204062_s_at	KIAA0623 gene product	Hs. 151406. 0	gb:BG526973		17	5. 3	286	2. 46	1. 68	0. 15
1193	Testis	204244_s_at	activator of S phase kinase	Hs. 152759. 0	gb:NM_006716. 1	ASK	7	5. 3	649	2. 81	1. 31	0. 28
1194	Testis	206526_at	DKFZP566F0546 protein	Hs. 144505. 0	gb:NM_015653. 1	DKFZP566F0	22	5. 3	164	2. 21	0. 59	0. 30
1195	Testis	212374_at	FEM-1 (C. elegans) homolog b	Hs. 6048. 0	gb:NM_015322. 1	FEM1B	15	5. 3	426	2. 63	1. 17	0. 27
1196	Testis	220749_at	hypothetical protein FLJ13031	Hs. 97757. 0	gb:NM_024688. 1	FLJ13031	10	5. 3	270	2. 43	1. 69	0. 14
1197	Testis	221845_s_at	suppressor of potassium transport defect 3	Hs. 21263. 1	gb:AI655698_RC		11	5. 3	626	2. 80	0. 77	0. 38
1198	Testis	206197_at	non-metastatic cells 5, protein expressed in (r	Hs. 72050. 0	gb:NM_003551. 1	NME5	5	5. 2	658	2. 82	1. 73	0. 21
1199	Testis	206318_at	eppin-3	Hs. 121084. 0	gb:NM_020398. 1	EPPIN3	20	5. 2	156	2. 19	1. 37	0. 16
1200	Testis	207021_at	zona pellucida binding protein	Hs. 99875. 0	gb:NM_007009. 1	SP38	7	5. 2	1463	3. 17	1. 38	0. 34
1201	Testis	207739_s_at	G antigen 2	Hs. 278444. 0	gb:NM_001472. 1	GAGE2	X	5. 2	808	2. 91	0. 88	0. 39
1202	Testis	213996_at	hypothetical protein	Hs. 71791. 0	gb:NM_013313. 1	AF060862	22	5. 2	912	2. 96	1. 43	0. 30
1203	Testis	214408_s_at	ret finger protein-like 3 antisense	Hs. 274285. 0	gb:AL021937		22	5. 2	205	2. 31	1. 32	0. 19
1204	Testis	205714_s_at	BLu protein	Hs. 167380. 0	gb:NM_015896. 1	BLu	3	5. 1	279	2. 45	0. 73	0. 34
1205	Testis	207213_s_at	ubiquitin specific protease 2	Hs. 125902. 0	gb:NM_004205. 1	USP2	11	5. 1	163	2. 21	0. 88	0. 26
1206	Testis	216778_s_at	cylicin, basic protein of sperm head cytoskelet	Hs. 307358. 0	gb:Z22780. 1			5. 1	234	2. 37	0. 97	0. 27
1207	Testis	219786_at	metallothionein-like 5, testis-specific (tesmir	Hs. 121378. 0	gb:NM_004923. 1	MTL5		5. 1	338	2. 53	1. 45	0. 21
1208	Testis	220766_at	B-cell translocation gene 4	Hs. 128180. 0	gb:NM_017589. 1	BTG4	11	5. 1	175	2. 24	0. 58	0. 32
1209	Testis	220790_s_at	membrane-spanning 4-domains, subfamily A, membe	Hs. 178066. 0	gb:NM_023945. 1	TETM4	11	5. 1	229	2. 36	0. 87	0. 29
1210	Testis	202170_s_at	aminoadipate-semialdehyde dehydrogenase-phospho	Hs. 64595. 0	gb:AF151057. 1		11	5	323	2. 51	1. 82	0. 14
1211	Testis	207144_s_at	Cbpp300-interacting transactivator, with GluAsp	Hs. 40403. 0	gb:NM_004143. 1	CITED1	X	5	597	2. 78	0. 94	0. 37
1212	Testis	207664_at	a disintegrin and metalloproteinase domain 2 (f	Hs. 177959. 0	gb:NM_001464. 2	ADAM2	8	5	222	2. 35	0. 47	0. 38
1213	Testis	210033_s_at	sperm associated antigen 6	Hs. 158213. 0	gb:AF079363. 1		10	5	286	2. 46	1. 50	0. 19
1214	Testis	210171_s_at	cAMP responsive element modulator	Hs. 155924. 2	gb:S68134. 1	CREM	10	5	270	2. 43	0. 30	0. 42
1215	Testis	210262_at	testis specific protein 1 (probe H4-1 p3-1)	Hs. 2042. 0	gb:M25532. 1	GAPDL5	6	5	2255	3. 35	0. 94	0. 49
1216	Testis	219199_at	ALL1 fused gene from 5q31	Hs. 231967. 0	gb:NM_014423. 1	AF5Q31	5	5	151	2. 18	1. 62	0. 11

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	A	B	C	D	E	F	G	H	I	J	K	L
1217	Testis	219703_at	hypothetical protein FLJ11222	Hs. 132851. 0	gb:NM_018365. 1	FLJ11222	15	5	504	2. 70	1. 19	0. 30
1218	Testis	207343_at	lysozyme homolog	Hs. 97477. 0	gb:NM_020426. 1	LOC57151	17	4. 9	230	2. 36	1. 73	0. 13
1219	Testis	209382_at	polymerase (RNA) III (DNA directed) (62kD)	Hs. 250745. 1	gb:U93867. 1	RPC62	1	4. 9	224	2. 35	1. 81	0. 11
1220	Testis	209923_s_at	BRCA1 associated protein	Hs. 122764. 0	gb:AF035620. 1	BRAP2	12	4. 9	227	2. 36	1. 06	0. 27
1221	Testis	211114_x_at	survival of motor neuron protein interacting pı	Hs. 102456. 2	gb:AB037702. 1	SIP1-gamma	14	4. 9	203	2. 31	1. 85	0. 09
1222	Testis	219148_at	PDZ-binding kinase; T-cell originated protein k	Hs. 104741. 0	gb:NM_018492. 1	TOPK	8	4. 9	644	2. 81	1. 34	0. 30
1223	Testis	219211_at	ubiquitin specific protease 18	Hs. 38260. 0	gb:NM_017414. 1	USP18	22	4. 9	161	2. 21	0. 93	0. 26
1224	Testis	219834_at	hypothetical protein FLJ21579	Hs. 122970. 0	gb:NM_024744. 1	FLJ21579	2	4. 9	98	1. 99	1. 21	0. 16
1225	Testis	219867_at	hypothetical protein FLJ12627	Hs. 283725. 0	gb:NM_024944. 1	FLJ12627	21	4. 9	199	2. 30	1. 10	0. 24
1226	Testis	221167_s_at			gb:NM_031290. 1	DKFZP434K1	13	4. 9	278	2. 44	1. 30	0. 24
1227	Testis	221252_s_at			gb:NM_031289. 1	MGC3146	12	4. 9	779	2. 89	1. 14	0. 36
1228	Testis	202046_s_at	glucocorticoid receptor DNA binding factor 1	Hs. 102548. 0	gb:NM_004491. 1	GRLF1	19	4. 8	212	2. 33	1. 49	0. 18
1229	Testis	203062_s_at	KIAA0170 gene product	Hs. 277585. 0	gb:NM_014641. 1	KIAA0170		4. 8	308	2. 49	1. 81	0. 14
1230	Testis	208365_s_at	G protein-coupled receptor kinase 2 (Drosophila	Hs. 32959. 0	gb:NM_005307. 1	GPRK2L		4. 8	447	2. 65	1. 36	0. 27
1231	Testis	210032_s_at	sperm associated antigen 6	Hs. 158213. 0	gb:AI651156_RC		10	4. 8	790	2. 90	1. 43	0. 31
1232	Testis	212380_at	KIAA0082 protein	Hs. 154045. 0	gb:D43949. 1	KIAA0082	6	4. 8	357	2. 55	1. 96	0. 12
1233	Testis	218077_s_at	DHHC1 protein	Hs. 14896. 0	gb:BE542551		3	4. 8	196	2. 29	1. 72	0. 12
1234	Testis	218327_s_at	synaptosomal-associated protein, 29kD	Hs. 194714. 0	gb:NM_004782. 1	SNAP29	22	4. 8	340	2. 53	1. 82	0. 15
1235	Testis	219870_at	hypothetical protein FLJ12668	Hs. 287540. 0	gb:NM_024997. 1	FLJ12668	16	4. 8	337	2. 53	1. 63	0. 19
1236	Testis	205899_at	cyclin A1	Hs. 79378. 0	gb:NM_003914. 1	CCNA1	13	4. 7	709	2. 85	1. 52	0. 28
1237	Testis	213186_at	KIAA0675 gene product	Hs. 165662. 1	gb:BG502305		3	4. 7	334	2. 52	1. 79	0. 16
1238	Testis	214868_at	piwi (Drosophila)-like 1	Hs. 194712. 0	gb:AW189518_RC		12	4. 7	360	2. 56	0. 87	0. 36
1239	Testis	215133_s_at	ribosomal protein L19	Hs. 252723. 1	gb:AL117630. 1		5	4. 7	352	2. 55	0. 71	0. 39
1240	Testis	216370_s_at	transketolase-like 1	Hs. 102866. 2	gb:Z49258			4. 7	625	2. 80	1. 36	0. 30
1241	Testis	218819_at	deleted in cancer 1; RNA helicase HDBDICE1	Hs. 58570. 0	gb:NM_012141. 1	DDX26	13	4. 7	310	2. 49	1. 85	0. 14
1242	Testis	220365_at	putative allantoicase	Hs. 97899. 0	gb:NM_018436. 1	LOC55821	2	4. 7	164	2. 21	0. 91	0. 28
1243	Testis	221316_at	hypothetical protein DKFZp434A1022	Hs. 324335. 0	gb:NM_021185. 1	DKFZP434A1	19	4. 7	290	2. 46	1. 09	0. 29
1244	Testis	207482_at	clg01 protein	Hs. 283640. 0	gb:NM_014477. 1	CLG01	20	4. 6	205	2. 31	1. 53	0. 17
1245	Testis	207663_x_at	G antigen 3	Hs. 176661. 0	gb:NM_001473. 1	GAGE3	X	4. 6	532	2. 73	1. 41	0. 29
1246	Testis	215044_s_at	Homo sapiens cDNA FLJ13428 fis, clone PLACE1002	Hs. 306650. 0	gb:AI571996_RC		2	4. 6	171	2. 23	1. 58	0. 14
1247	Testis	217250_s_at	KIAA0444 protein	Hs. 158291. 1	gb:AL110281. 1		1	4. 6	341	2. 53	1. 28	0. 27
1248	Testis	219785_s_at	hypothetical protein FLJ22477	Hs. 118944. 0	gb:NM_024735. 1	FLJ22477	16	4. 6	518	2. 71	1. 25	0. 32
1249	Testis	221673_s_at	casein kinase 1, gamma 1	Hs. 157777. 1	gb:AB042563. 1	CSNK1G1L		4. 6	187	2. 27	1. 45	0. 18
1250	Testis	222253_s_at	DKFZp434P211 protein	Hs. 278387. 1	gb:AL117484. 1		22	4. 6	570	2. 76	1. 14	0. 35
1251	Testis	37425_g_at						4. 5	161	2. 21	1. 30	0. 20
1252	Testis	50400_at						4. 5	289	2. 46	1. 54	0. 21
1253	Testis	203176_s_at	transcription factor 6-like 1 (mitochondrial tı	Hs. 75133. 0	gb:BE552470_RC		10	4. 5	225	2. 35	1. 76	0. 13
1254	Testis	206172_at	interleukin 13 receptor, alpha 2	Hs. 25954. 0	gb:NM_000640. 1	IL13RA2	X	4. 5	491	2. 69	1. 04	0. 37

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	A	B	C	D	E	F	G	H	I	J	K	L
1255	Testis	210235_s_at	protein tyrosine phosphatase, receptor type, f	Hs. 183648. 1	gb:U22815. 1		11	4. 5	224	2. 35	1. 70	0. 14
1256	Testis	210600_s_at	G protein-coupled receptor kinase 2 (Drosophil	Hs. 32959. 1	gb:U33054. 1		4	4. 5	148	2. 17	0. 18	0. 44
1257	Testis	214392_at	hypothetical protein R30953_1	Hs. 180943. 0	gb:AA431984_RC		19	4. 5	110	2. 04	1. 25	0. 18
1258	Testis	214838_at	H. sapiens gene from PAC 747L4	Hs. 119254. 0	gb:AL035297. 1		1	4. 5	269	2. 43	1. 96	0. 10
1259	Testis	215486_at	Homo sapiens phosphoribosylpyrophosphate synth	Hs. 169284. 0	gb:AW072461_RC		7	4. 5	336	2. 53	1. 18	0. 30
1260	Testis	218695_at	exosome component Rrp41	Hs. 97574. 0	gb:NM_019037. 1	FLJ20591		4. 5	183	2. 26	1. 16	0. 24
1261	Testis	219945_at	gonadotropin-regulated testicular RNA helicase	Hs. 98738. 0	gb:NM_013264. 1	GRTH	11	4. 5	550	2. 74	0. 90	0. 41
1262	Testis	220390_at	hypothetical protein FLJ23598	Hs. 147377. 0	gb:NM_024783. 1	FLJ23598	11	4. 5	127	2. 10	1. 33	0. 17
1263	Testis	222075_s_at	Homo sapiens ornithine decarboxylase antizyme	Hs. 326705. 0	gb:BE792072		1	4. 5	1069	3. 03	1. 24	0. 40
1264	Testis	222158_s_at	CGI-146 protein	Hs. 42409. 4	gb:AF229834. 1	PNAS-4	1	4. 5	289	2. 46	1. 72	0. 16
1265	Testis	203654_s_at	coilin	Hs. 966. 0	gb:NM_004645. 1	COIL	17	4. 4	657	2. 82	1. 72	0. 25
1266	Testis	205282_at	low density lipoprotein receptor-related protei	Hs. 54481. 0	gb:NM_004631. 1	LRP8	1	4. 4	212	2. 33	1. 51	0. 19
1267	Testis	206550_s_at	nucleoporin 155kD	Hs. 23255. 0	gb:NM_004298. 1	NUP155	5	4. 4	365	2. 56	0. 99	0. 36
1268	Testis	206985_at	hydroxysteroid (17-beta) dehydrogenase 3	Hs. 477. 0	gb:NM_000197. 1	HSD17B3	9	4. 4	354	2. 55	1. 70	0. 19
1269	Testis	207405_s_at	RAD17 (S. pombe) homolog	Hs. 16184. 0	gb:NM_002873. 1	RAD17	5	4. 4	286	2. 46	1. 60	0. 20
1270	Testis	210323_at	tektin 2 (testicular)	Hs. 127111. 0	gb:AB033823. 1	h-tektin-t	1	4. 4	187	2. 27	0. 86	0. 32
1271	Testis	211381_x_at	sperm associated antigen 11	Hs. 2717. 2	gb:AF168617. 1		8	4. 4	324	2. 51	0. 94	0. 35
1272	Testis	214296_x_at	Homo sapiens chromosome 19, cosmid R26660	Hs. 90010. 0	gb:AV721013		NA	4. 4	240	2. 38	1. 02	0. 31
1273	Testis	219721_at	hypothetical protein FLJ11181	Hs. 28472. 0	gb:NM_018350. 1	FLJ11181	15	4. 4	323	2. 51	0. 71	0. 40
1274	Testis	219904_at	hypothetical protein MGC4161	Hs. 177688. 0	gb:NM_024303. 1	MGC4161	19	4. 4	149	2. 17	1. 60	0. 13
1275	Testis	221446_at	a disintegrin and metalloproteinase domain 30	Hs. 283011. 0	gb:NM_021794. 1	ADAM30		4. 4	238	2. 38	1. 39	0. 22
1276	Testis	59631_at						4. 3	262	2. 42	1. 59	0. 19
1277	Testis	202631_s_at	amyloid beta precursor protein (cytoplasmic ta	Hs. 84084. 0	gb:NM_006380. 1	APPBP2	17	4. 3	227	2. 36	1. 59	0. 18
1278	Testis	203664_s_at	polymerase (RNA) II (DNA directed) polypeptide	Hs. 194638. 0	gb:NM_004805. 1	POLR2D	2	4. 3	353	2. 55	1. 76	0. 18
1279	Testis	204325_s_at	neurofibromin 1 (neurofibromatosis, von Recklin	Hs. 93207. 0	gb:NM_000267. 1	NF1	17	4. 3	148	2. 17	1. 66	0. 12
1280	Testis	205034_at	cyclin E2	Hs. 30464. 0	gb:NM_004702. 1	CCNE2	8	4. 3	136	2. 13	1. 26	0. 20
1281	Testis	206588_at	deleted in azoospermia-like	Hs. 73078. 0	gb:NM_001351. 1	DAZL	3	4. 3	491	2. 69	1. 06	0. 38
1282	Testis	208132_x_at	HLA-B associated transcript-2	Hs. 25911. 0	gb:NM_004638. 1	D6S51E	6	4. 3	351	2. 54	1. 94	0. 14
1283	Testis	213599_at	Opa-interacting protein 5	Hs. 116206. 0	gb:BE045993_RC		15	4. 3	261	2. 42	1. 60	0. 19
1284	Testis	218219_s_at	TASP for testis-specific adriamycin sensitivity	Hs. 134342. 0	gb:NM_018697. 1	LOC55915	7	4. 3	212	2. 33	1. 63	0. 16
1285	Testis	218305_at	hypothetical protein FLJ23338	Hs. 61790. 0	gb:NM_024658. 1	FLJ23338	14	4. 3	268	2. 43	1. 71	0. 17
1286	Testis	219753_at	stromal antigen 3	Hs. 20132. 0	gb:NM_012447. 1	STAG3	7	4. 3	634	2. 80	1. 36	0. 34
1287	Testis	220545_s_at	testis-specific kinase substrate	Hs. 224210. 0	gb:NM_021733. 1	TSKS	19	4. 3	209	2. 32	1. 55	0. 18
1288	Testis	203702_s_at	KIAA0173 gene product	Hs. 169910. 0	gb:AL043927_RC		2	4. 2	179	2. 25	1. 07	0. 28
1289	Testis	208307_at	RNA binding motif protein, Y chromosome, famil	Hs. 2958. 0	gb:NM_005058. 1	RBMY1A1	Y	4. 2	163	2. 21	0. 97	0. 29
1290	Testis	215929_at	Homo sapiens mRNA; cDNA DKFZp434F083 (from clo	Hs. 119207. 0	gb:AL080132. 1		10	4. 2	214	2. 33	0. 70	0. 39
1291	Testis	220190_s_at	stoned BTFIIA-alphabeta-like factor	Hs. 54961. 0	gb:NM_006873. 1	SALF	2	4. 2	203	2. 31	0. 32	0. 48
1292	Testis	220217_x_at	SPANX family, member C	Hs. 178053. 0	gb:NM_022661. 1	SPANXC	X	4. 2	376	2. 58	1. 49	0. 26

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	A	B	C	D	E	F	G	H	I	J	K	L
1293	Testis	205457_at	hypothetical protein MGC4614	Hs. 300691. 0	gb:NM_024294. 1	MGC4614	6	4. 1	281	2. 45	1. 50	0. 23
1294	Testis	206721_at	hypothetical protein LOC57821	Hs. 122547. 0	gb:NM_021179. 1	LOC57821	1	4. 1	131	2. 12	1. 11	0. 25
1295	Testis	207209_at	centrin, EF-hand protein, 1	Hs. 122511. 0	gb:NM_004066. 1	CETN1	18	4. 1	333	2. 52	1. 74	0. 19
1296	Testis	210534_s_at	B9 protein	Hs. 11955. 1	gb:BC002944. 1		17	4. 1	252	2. 40	1. 73	0. 16
1297	Testis	213761_at	hypothetical protein	Hs. 12871. 0	gb:AW664850_RC		12	4. 1	184	2. 27	1. 61	0. 16
1298	Testis	214659_x_at	ZAP3 protein	Hs. 159471. 3	gb:AC007956		14	4. 1	162	2. 21	1. 69	0. 13
1299	Testis	217099_s_at	gemin4	Hs. 19614. 1	gb:AF258545			4. 1	323	2. 51	1. 86	0. 16
1300	Testis	218737_at	hypothetical protein FLJ10701	Hs. 146589. 0	gb:NM_018183. 1	FLJ10701	12	4. 1	99	1. 99	1. 38	0. 15
1301	Testis	221941_at	ESTs, Weakly similar to T47142 hypothetical pr	Hs. 292503. 0	gb:BE646315_RC		10	4. 1	203	2. 31	1. 69	0. 15
1302	Testis	222089_s_at	KIAA0557 protein	Hs. 101414. 0	gb:BF508774_RC		16	4. 1	182	2. 26	0. 85	0. 34
1303	Testis	222096_x_at	Homo sapiens cDNA FLJ12825 fis, clone NT2RP200	Hs. 300683. 0	gb:BF056462_RC		9	4. 1	171	2. 23	1. 11	0. 27
1304	Testis	57703_at						4	93	1. 97	1. 42	0. 14
1305	Testis	204233_s_at	choline kinase	Hs. 77221. 0	gb:AI991328_RC		11	4	499	2. 70	1. 52	0. 29
1306	Testis	205673_s_at	DKFZP564L0862 protein	Hs. 19404. 0	gb:NM_024087. 1	DKFZP564L0	X	4	159	2. 20	1. 41	0. 20
1307	Testis	207281_x_at	variable charge protein on X with two repeats	Hs. 279737. 0	gb:NM_016378. 1	VCX-2r	X	4	283	2. 45	0. 97	0. 37
1308	Testis	207427_at	acrosin	Hs. 183088. 0	gb:NM_001097. 1	ACR	22	4	439	2. 64	1. 21	0. 36
1309	Testis	213392_at	G protein-coupled receptor, family C, group 5,	Hs. 242407. 2	gb:AW070229_RC		16	4	404	2. 61	1. 86	0. 19
1310	Testis	218370_s_at	hypothetical protein FLJ12903	Hs. 14928. 0	gb:NM_022753. 1	FLJ12903	1	4	262	2. 42	1. 78	0. 16
1311	Testis	220141_at	hypothetical protein FLJ23554	Hs. 164705. 0	gb:NM_024806. 1	FLJ23554	11	4	280	2. 45	1. 37	0. 27
1312	Testis	220493_at	doublesex and mab-3 related transcription fact	Hs. 98586. 0	gb:NM_021951. 1	DMRT1	9	4	238	2. 38	0. 90	0. 37
1313	Testis	221849_s_at	Homo sapiens, clone IMAGE:3953631, mRNA, parti	Hs. 298553. 0	gb:AA733079_RC		19	4	82	1. 91	0. 88	0. 26
1314	Testis	206440_at	Vertebrate LIN7 homolog 1, Tax interaction pro	Hs. 178215. 0	gb:NM_004664. 1	VELI1	12	3. 9	135	2. 13	0. 76	0. 35
1315	Testis	206787_at	bromodomain, testis-specific	Hs. 178281. 0	gb:NM_001726. 1	BRDT	1	3. 9	268	2. 43	0. 66	0. 46
1316	Testis	210437_at	melanoma antigen, family A, 9	Hs. 37110. 0	gb:BC002351. 1		X	3. 9	121	2. 08	0. 84	0. 32
1317	Testis	210465_s_at	small nuclear RNA activating complex, polypepti	Hs. 164915. 1	gb:U71300. 1	SNAP50	9	3. 9	149	2. 17	1. 66	0. 13
1318	Testis	211238_at	a disintegrin and metalloproteinase domain 7	Hs. 166003. 0	gb:AF215824. 1	ADAM7	8	3. 9	124	2. 09	0. 56	0. 39
1319	Testis	218628_at	CGI-116 protein	Hs. 18885. 0	gb:NM_016053. 1	LOC51019	12	3. 9	489	2. 69	1. 67	0. 26
1320	Testis	219984_s_at	H-REV107 protein-related protein	Hs. 36761. 0	gb:NM_020386. 1	LOC57110	3	3. 9	505	2. 70	1. 16	0. 40
1321	Testis	220618_s_at	hypothetical protein FLJ10057	Hs. 105191. 0	gb:NM_017984. 1	FLJ10057	7	3. 9	224	2. 35	1. 19	0. 30
1322	Testis	222157_s_at	KIAA1449 protein	Hs. 109778. 1	gb:AB040882. 1	KIAA1449	3	3. 9	296	2. 47	1. 88	0. 15
1323	Testis	37424_at						3. 8	300	2. 48	1. 73	0. 19
1324	Testis	204353_s_at	DKFZP586D211 protein	Hs. 31968. 0	gb:BC002923. 1		7	3. 8	192	2. 28	1. 74	0. 14
1325	Testis	205443_at	small nuclear RNA activating complex, polypepti	Hs. 179312. 0	gb:NM_003082. 1	SNAPC1	14	3. 8	134	2. 13	1. 47	0. 17
1326	Testis	205514_at	hypothetical protein FLJ11191	Hs. 21838. 0	gb:NM_018355. 1	FLJ11191	19	3. 8	112	2. 05	1. 38	0. 18
1327	Testis	207678_s_at	SRY (sex determining region Y)-box 30	Hs. 197805. 0	gb:NM_007017. 1	SOX30	5	3. 8	130	2. 12	0. 55	0. 41
1328	Testis	210843_s_at	KIAA0626 gene product	Hs. 178121. 1	gb:AF327560. 1			3. 8	200	2. 30	1. 43	0. 23
1329	Testis	213956_at	KIAA0480 gene product	Hs. 92200. 1	gb:AW299294_RC		1	3. 8	229	2. 36	1. 67	0. 18
1330	Testis	214222_at	KIAA0944 protein	Hs. 97403. 0	gb:AB023161. 1	KIAA0944	2	3. 8	147	2. 17	1. 40	0. 20

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	A	B	C	D	E	F	G	H	I	J	K	L
1331	Testis	217004_s_at	MCF.2 cell line derived transforming sequence	Hs.89543.3	gb:X13230.1		X	3.8	105	2.02	1.26	0.20
1332	Testis	217380_s_at	Homo sapiens clone DT1P1A11 mRNA, CAG repeat re	Hs.168303.0	gb:U92992.1			3.8	149	2.17	0.58	0.42
1333	Testis	220058_at	hypothetical protein MGC3048	Hs.115437.0	gb:NM_024052.1	MGC3048		3.8	377	2.58	1.07	0.39
1334	Testis	220991_s_at			gb:NM_030936.1	DKFZP434C1	7	3.8	222	2.35	0.57	0.47
1335	Testis	222327_x_at	olfactory receptor, family 7, subfamily E, memb	Hs.306191.0	gb:AA459867_RC		13	3.8	551	2.74	1.89	0.23
1336	Testis	202924_s_at	pleiomorphic adenoma gene-like 2	Hs.154104.0	gb:AL562280_RC		20	3.7	246	2.39	1.71	0.18
1337	Testis	203956_at	KIAA0852 protein	Hs.35276.0	gb:NM_014941.1	KIAA0852	22	3.7	237	2.38	1.78	0.16
1338	Testis	205527_s_at	DKFZP434D174 protein	Hs.302421.0	gb:NM_015487.1	DKFZP434D1	17	3.7	198	2.30	0.96	0.36
1339	Testis	211080_s_at			gb:Z25425.1		1	3.7	176	2.25	1.26	0.27
1340	Testis	213410_at	DKFZP586F1019 protein	Hs.227209.0	gb:AL050102.1	DKFZp586F1	10	3.7	250	2.40	1.86	0.15
1341	Testis	215472_at	Human DNA sequence from clone RP1-257A15 on ch	Hs.144339.0	gb:AI017119_RC		6	3.7	209	2.32	1.46	0.23
1342	Testis	217094_s_at	atrophin-1 interacting protein 4	Hs.98074.1	gb:AL109923			3.7	259	2.41	1.45	0.26
1343	Testis	217124_at	KIAA1023 protein	Hs.21361.1	gb:AL136792.1	DKFZp434P1219		3.7	250	2.40	1.19	0.33
1344	Testis	219311_at	hypothetical protein FLJ12542	Hs.236940.0	gb:NM_024899.1	FLJ12542	18	3.7	160	2.21	0.83	0.38
1345	Testis	220230_s_at	cytochrome b5 reductase b5R.2	Hs.22142.0	gb:NM_016229.1	LOC51700		3.7	592	2.77	1.67	0.30
1346	Testis	221048_x_at	hypothetical protein FLJ20721	Hs.12929.0	gb:NM_017941.1	FLJ20721	17	3.7	98	1.99	1.05	0.25
1347	Testis	221078_s_at	hypothetical protein FLJ10392	Hs.20887.0	gb:NM_018084.1	FLJ10392	2	3.7	172	2.24	1.18	0.29
1348	Testis	201504_s_at	translin	Hs.75066.0	gb:AI435302_RC		2	3.6	147	2.17	1.14	0.29
1349	Testis	204250_s_at	KIAA1052 protein	Hs.18624.0	gb:AI655714_RC		11	3.6	103	2.01	1.23	0.22
1350	Testis	204377_s_at	KIAA0800 gene product	Hs.118738.0	gb:NM_014703.1	KIAA0800	3	3.6	344	2.54	1.39	0.31
1351	Testis	205186_at	dynein, axonemal, light intermediate polypeptid	Hs.33846.0	gb:NM_003462.2	P28	1	3.6	325	2.51	1.13	0.38
1352	Testis	205588_s_at	FGFR1 oncogene partner	Hs.180296.0	gb:NM_007045.1	FOP	6	3.6	384	2.58	1.43	0.32
1353	Testis	206819_at	DKFZp434P211 protein	Hs.278387.0	gb:NM_014549.1	DKFZP434P2	22	3.6	295	2.47	1.26	0.34
1354	Testis	209942_x_at	melanoma antigen, family A, 3	Hs.36978.0	gb:BC000340.1		X	3.6	229	2.36	1.27	0.30
1355	Testis	210189_at	heat shock 70kD protein-like 1	Hs.80288.0	gb:D85730.1	HSPA1L	6	3.6	453	2.66	1.15	0.41
1356	Testis	214015_at	Nck, Ash and phospholipase C binding protein	Hs.104481.0	gb:AI968839_RC		17	3.6	328	2.52	1.20	0.37
1357	Testis	219104_at	C3HC4-like zinc finger protein	Hs.44685.0	gb:NM_016422.1	ZFP26	11	3.6	177	2.25	1.61	0.18
1358	Testis	220051_at	protease, serine, 21 (testisin)	Hs.72026.0	gb:NM_006799.1	PRSS21	16	3.6	416	2.62	1.56	0.29
1359	Testis	220129_at	hypothetical protein FLJ20449	Hs.130546.0	gb:NM_017826.1	FLJ20449	13	3.6	148	2.17	1.23	0.26
1360	Testis	220278_at	hypothetical protein FLJ10251	Hs.120842.0	gb:NM_018039.1	FLJ10251	11	3.6	257	2.41	1.02	0.38
1361	Testis	221998_s_at	VRK3 for vaccinia related kinase 3	Hs.98289.1	gb:BF062886_RC		19	3.6	536	2.73	1.75	0.27
1362	Testis	222224_at	Homo sapiens alpha-NAC gene for nascent polypep	Hs.287764.0	gb:AJ278883			3.6	73	1.87	0.83	0.29
1363	Testis	203730_s_at	zinc finger protein homologous to Zfp95 in mous	Hs.110839.0	gb:BF196931_RC		7	3.5	123	2.09	1.54	0.16
1364	Testis	204641_at	NIMA (never in mitosis gene a)-related kinase 2	Hs.153704.0	gb:NM_002497.1	NEK2	1	3.5	580	2.76	0.77	0.58
1365	Testis	204843_s_at	protein kinase, cAMP-dependent, regulatory, typ	Hs.289075.0	gb:NM_004157.1	PRKAR2A	3	3.5	241	2.38	1.54	0.24
1366	Testis	206547_s_at	protein phosphatase, EF hand calcium-binding do	Hs.211589.0	gb:NM_006240.1	PPEF1	X	3.5	130	2.11	1.32	0.22
1367	Testis	211702_s_at			gb:AF350251.1		17	3.5	202	2.30	1.70	0.17
1368	Testis	212949_at	KIAA0074 protein	Hs.1192.0	gb:D38553.1	HCAP-H	2	3.5	194	2.29	1.09	0.34

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	A	B	C	D	E	F	G	H	I	J	K	L
1369	Testis	213644_at	ESTs, Weakly similar to KIAA1276 protein H. sapi	Hs. 177482. 0	gb:AI979276_RC		17	3. 5	452	2. 65	1. 41	0. 36
1370	Testis	214431_at	guanine monphosphate synthetase	Hs. 5398. 0	gb:NM_003875. 1	GMPS	3	3. 5	346	2. 54	1. 90	0. 18
1371	Testis	214706_at	zinc finger protein 200	Hs. 88219. 1	gb:AU149447_RC		16	3. 5	157	2. 19	1. 64	0. 16
1372	Testis	215461_at	Homo sapiens chromosome 19, cosmid R31343	Hs. 126496. 0	gb:AI807507_RC		19	3. 5	328	2. 52	1. 36	0. 33
1373	Testis	216917_s_at	synaptonemal complex protein 1	Hs. 112743. 1	gb:X95654. 1	SCP1	1	3. 5	205	2. 31	0. 94	0. 39
1374	Testis	217520_x_at	ESTs, Moderately similar to HERC2 H. sapiens	Hs. 154999. 0	gb:BG396614		15	3. 5	228	2. 36	1. 70	0. 19
1375	Testis	219258_at	hypothetical protein FLJ20516	Hs. 70811. 0	gb:NM_017858. 1	FLJ20516	15	3. 5	99	1. 99	0. 80	0. 34
1376	Testis	220004_at	DEAD-box protein	Hs. 125507. 0	gb:NM_018665. 1	HAGE	6	3. 5	100	2. 00	0. 48	0. 43
1377	Testis	220325_at	hypothetical protein FLJ23157	Hs. 223806. 0	gb:NM_024885. 1	TAF2Q	X	3. 5	190	2. 28	1. 25	0. 29
1378	Testis	222149_x_at	Homo sapiens mRNA; cDNA DKFZp434K052 (from clor	Hs. 169639. 0	gb:AL137398. 1		15	3. 5	195	2. 29	1. 39	0. 26
1379	Testis	51146_at						3. 4	143	2. 16	1. 41	0. 22
1380	Testis	204782_at	myeloid leukemia factor 1	Hs. 85195. 0	gb:NM_022443. 1_I	MLF1	3	3. 4	86	1. 93	0. 65	0. 37
1381	Testis	205224_at	surfeit 2	Hs. 159448. 0	gb:NM_017503. 1	SURF2	9	3. 4	88	1. 95	0. 95	0. 29
1382	Testis	207523_at	chromosome 6 open reading frame 10	Hs. 57692. 0	gb:NM_006781. 1	TSBP	6	3. 4	156	2. 19	0. 89	0. 38
1383	Testis	209799_at	protein kinase, AMP-activated, alpha 1 catalyti	Hs. 9247. 0	gb:AF100763. 1		5	3. 4	276	2. 44	1. 32	0. 33
1384	Testis	217559_at	ESTs, Highly similar to A42735 ribosomal protei	Hs. 308332. 0	gb:AI001784_RC		14	3. 4	288	2. 46	1. 26	0. 36
1385	Testis	218480_at	hypothetical protein FLJ21839	Hs. 285122. 0	gb:NM_021831. 1	FLJ21839	2	3. 4	559	2. 75	1. 38	0. 40
1386	Testis	218993_at	hypothetical protein FLJ10581	Hs. 182729. 0	gb:NM_018146. 1	FLJ10581	17	3. 4	254	2. 40	1. 89	0. 15
1387	Testis	219583_s_at	hypothetical protein	Hs. 177011. 0	gb:NM_018418. 1	HSD-3. 1	14	3. 4	244	2. 39	1. 21	0. 34
1388	Testis	220269_at	hypothetical protein FLJ23049	Hs. 96901. 0	gb:NM_024687. 1	FLJ23049		3. 4	250	2. 40	0. 93	0. 43
1389	Testis	220382_s_at	hypothetical protein FLJ10312	Hs. 132560. 0	gb:NM_030672. 1	FLJ10312	18	3. 4	269	2. 43	1. 05	0. 41
1390	Testis	220623_s_at	testis specific, 10	Hs. 116116. 0	gb:NM_025244. 1	TSGA10	2	3. 4	305	2. 48	1. 20	0. 37
1391	Testis	221973_at	fibulin 1	Hs. 79732. 5	gb:AI983904_RC		2	3. 4	400	2. 60	1. 39	0. 36
1392	Testis	42361_g_at						3. 3	158	2. 20	1. 28	0. 27
1393	Testis	202044_at	glucocorticoid receptor DNA binding factor 1	Hs. 102548. 0	gb:AU159484_RC		19	3. 3	180	2. 26	1. 25	0. 30
1394	Testis	203181_x_at	SFRS protein kinase 2	Hs. 78353. 0	gb:AW149364_RC		7	3. 3	289	2. 46	1. 54	0. 28
1395	Testis	206412_at	fer (fpsfes related) tyrosine kinase (phosphopr	Hs. 121558. 0	gb:NM_005246. 1	FER	5	3. 3	61	1. 79	0. 89	0. 27
1396	Testis	206683_at	zinc finger protein 165	Hs. 55481. 0	gb:NM_003447. 1	ZNF165	6	3. 3	182	2. 26	0. 49	0. 53
1397	Testis	207597_at	a disintegrin and metalloproteinase domain 18	Hs. 127930. 0	gb:NM_014237. 1	ADAM18	8	3. 3	133	2. 12	1. 35	0. 23
1398	Testis	210115_at	Homo sapiens ribosomal protein L39 mRNA, comple	Hs. 132748. 0	gb:L05096. 1		3	3. 3	314	2. 50	0. 96	0. 46
1399	Testis	210394_x_at	synovial sarcoma, X breakpoint 2	Hs. 289105. 3	gb:BC005325. 1		X	3. 3	173	2. 24	1. 48	0. 23
1400	Testis	213959_s_at	KIAA1005 protein	Hs. 12328. 0	gb:BF515597_RC		16	3. 3	93	1. 97	1. 08	0. 27
1401	Testis	214229_at	dynein, axonemal, heavy polypeptide 17	Hs. 144672. 0	gb:AL122077. 1	DKFZp434H2	17	3. 3	156	2. 19	1. 51	0. 21
1402	Testis	218781_at	hypothetical protein FLJ22116	Hs. 34497. 0	gb:NM_024624. 1	FLJ22116	2	3. 3	324	2. 51	1. 78	0. 22
1403	Testis	219078_at	hypothetical protein FLJ10252	Hs. 53913. 0	gb:NM_018040. 1	FLJ10252	1	3. 3	251	2. 40	1. 71	0. 21
1404	Testis	220168_at	hypothetical protein FLJ10921	Hs. 279012. 0	gb:NM_018272. 1	FLJ10921	12	3. 3	153	2. 18	0. 86	0. 40
1405	Testis	220840_s_at	hypothetical protein FLJ10706	Hs. 273193. 0	gb:NM_018186. 1	FLJ10706	1	3. 3	74	1. 87	0. 76	0. 34
1406	Testis	220921_at	sperm protein associated with the nucleus, X cl	Hs. 293266. 0	gb:NM_013453. 1	SPANXA1	X	3. 3	103	2. 01	0. 41	0. 48

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	A	B	C	D	E	F	G	H	I	J	K	L
1407	Testis	202188_at	KIAA0095 gene product	Hs. 155314. 0	gb:NM_014669. 1	KIAA0095	16	3. 2	136	2. 13	1. 18	0. 30
1408	Testis	206483_at	testis specific leucine rich repeat protein	Hs. 57693. 0	gb:NM_012472. 1	TSLRP	8	3. 2	230	2. 36	1. 41	0. 30
1409	Testis	207617_at	DEADH (Asp-Glu-Ala-AspHis) box polypeptide 3	Hs. 147916. 1	gb:NM_024005. 1	DDX3	X	3. 2	92	1. 96	0. 79	0. 36
1410	Testis	210129_s_at	DKFZP434B103 protein	Hs. 289010. 1	gb:AF078842. 1		3	3. 2	195	2. 29	1. 33	0. 30
1411	Testis	212954_at	dual-specificity tyrosine-(Y)-phosphorylation	Hs. 17154. 0	gb:AF263541. 1	DYRK4	12	3. 2	258	2. 41	1. 52	0. 28
1412	Testis	214396_s_at	methyl-CpG binding domain protein 2	Hs. 25674. 2	gb:AI827820_RC		18	3. 2	234	2. 37	1. 14	0. 38
1413	Testis	214766_s_at	DKFZP434N093 protein	Hs. 33363. 1	gb:AL080144. 1	DKFZp434N0	2	3. 2	185	2. 27	1. 49	0. 24
1414	Testis	218123_at	chromosome 21 open reading frame 59	Hs. 5811. 0	gb:NM_017835. 1	C21ORF59	21	3. 2	252	2. 40	1. 79	0. 19
1415	Testis	220125_at	dynein, axonemal, intermediate polypeptide, 1	Hs. 112667. 0	gb:NM_012144. 1	DNAI1	9	3. 2	191	2. 28	1. 21	0. 34
1416	Testis	220238_s_at	SBBI26 protein	Hs. 26481. 0	gb:NM_018846. 1	SBBI26	7	3. 2	154	2. 19	0. 99	0. 37
1417	Testis	221992_at	KIAA0220 protein	Hs. 110613. 0	gb:AI925734_RC		16	3. 2	195	2. 29	1. 59	0. 22
1418	Testis	203086_at	kinesin heavy chain member 2	Hs. 113319. 0	gb:BE872563		5	3. 1	190	2. 28	1. 62	0. 21
1419	Testis	203350_at	adaptor-related protein complex 1, gamma 1 sub	Hs. 5344. 0	gb:NM_001128. 1	AP1G1	16	3. 1	233	2. 37	1. 70	0. 22
1420	Testis	203653_s_at	coilin	Hs. 966. 0	gb:BG391060		17	3. 1	225	2. 35	1. 15	0. 39
1421	Testis	206546_at	synaptonemal complex protein 2	Hs. 202676. 0	gb:NM_014258. 1	SYCP2	20	3. 1	125	2. 10	0. 86	0. 39
1422	Testis	207754_at	chromosome 12 open reading frame 2	Hs. 28529. 0	gb:NM_007211. 1	HOJ-1	12	3. 1	82	1. 91	1. 01	0. 29
1423	Testis	207962_at	calpain 11	Hs. 225953. 0	gb:NM_007058. 1	CAPN11		3. 1	100	2. 00	1. 01	0. 32
1424	Testis	209715_at	chromobox homolog 5 (Drosophila HP1 alpha)	Hs. 89232. 0	gb:L07515. 1		12	3. 1	132	2. 12	1. 38	0. 24
1425	Testis	214381_at	CDC10 (cell division cycle 10, S. cerevisiae,	Hs. 184326. 2	gb:BG150564_RC		9	3. 1	240	2. 38	1. 43	0. 31
1426	Testis	214612_x_at	melanoma antigen, family A, 6	Hs. 278460. 0	gb:U10691		X	3. 1	144	2. 16	1. 03	0. 36
1427	Testis	218296_x_at	hypothetical protein FLJ10504	Hs. 279763. 0	gb:NM_018116. 1	FLJ10504	1	3. 1	233	2. 37	1. 74	0. 20
1428	Testis	202666_s_at	BAF53	Hs. 274350. 0	gb:NM_004301. 1	BAF53A	3	3	183	2. 26	1. 73	0. 18
1429	Testis	202900_s_at	nucleoporin 88kD	Hs. 172108. 0	gb:NM_002532. 2	NUP88	17	3	341	2. 53	1. 85	0. 22
1430	Testis	203155_at	SET domain, bifurcated 1	Hs. 20991. 0	gb:NM_012432. 1	SETDB1	1	3	271	2. 43	1. 82	0. 20
1431	Testis	204157_s_at	KIAA0999 protein	Hs. 4278. 0	gb:NM_025164. 1	KIAA0999	11	3	425	2. 63	1. 81	0. 27
1432	Testis	204383_at	DiGeorge syndrome critical region gene DGSI	Hs. 154879. 0	gb:NM_022719. 1	DGSI	22	3	170	2. 23	1. 66	0. 19
1433	Testis	207374_at	phospholipid scramblase 2	Hs. 123411. 0	gb:NM_020359. 1	PLSCR2		3	154	2. 19	1. 31	0. 29
1434	Testis	222384_at	ESTs	Hs. 193769. 0	gb:AA417256_RC		1	3	55	1. 74	0. 76	0. 33
1435	Testis	205429_s_at	MAGUK protein p55T; Protein Associated with Lin	Hs. 108931. 0	gb:NM_016447. 1	LOC51678	7	2. 9	127	2. 10	0. 78	0. 45
1436	Testis	205662_at	B9 protein	Hs. 11955. 0	gb:NM_015681. 1	B9	17	2. 9	119	2. 07	1. 13	0. 33
1437	Testis	207228_at	protein kinase, cAMP-dependent, catalytic, gamm	Hs. 158029. 0	gb:NM_002732. 1	PRKACG	9	2. 9	119	2. 08	1. 18	0. 31
1438	Testis	210118_s_at	interleukin 1, alpha	Hs. 1722. 1	gb:M15329. 1	IL1A	2	2. 9	121	2. 08	1. 24	0. 30
1439	Testis	219715_s_at	hypothetical protein FLJ11090	Hs. 238039. 0	gb:NM_018319. 1	FLJ11090	14	2. 9	323	2. 51	1. 82	0. 24
1440	Testis	207056_s_at	solute carrier family 4, sodium bicarbonate co	Hs. 132136. 0	gb:NM_004858. 1	SLC4A8	12	2. 8	87	1. 94	0. 66	0. 46
1441	Testis	207666_x_at	synovial sarcoma, X breakpoint 3	Hs. 178749. 0	gb:NM_021014. 1	SSX3	X	2. 8	177	2. 25	1. 17	0. 38
1442	Testis	211865_s_at	Fzr1 protein	Hs. 268384. 1	gb:AB013463. 1	fzr2	19	2. 8	126	2. 10	1. 03	0. 38
1443	Testis	220544_at	testis-specific kinase substrate	Hs. 224210. 0	gb:NM_021733. 1	TSKS	19	2. 8	119	2. 07	1. 43	0. 23
1444	Testis	204266_s_at	choline kinase	Hs. 77221. 0	gb:NM_001277. 1	CHK	11	2. 7	495	2. 69	1. 80	0. 33

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	A	B	C	D	E	F	G	H	I	J	K	L
1445	Testis	216276_s_at	a disintegrin and metalloproteinase domain 3a	Hs. 98848. 0	gb:X89657. 1		16	2. 7	55	1. 74	0. 56	0. 43
1446	Testis	220381_at	hypothetical protein FLJ10312	Hs. 132560. 0	gb:NM_030672. 1	FLJ10312	18	2. 7	89	1. 95	1. 26	0. 25
1447	Testis	207580_at	melanoma antigen, family B, 4	Hs. 113291. 0	gb:NM_002367. 1	MAGEB4	X	2. 6	78	1. 89	0. 69	0. 47
1448	Testis	210536_s_at	sperm adhesion molecule 1 (PH-20 hyaluronidase,	Hs. 121494. 0	gb:S67798. 1	SPAM1	7	2. 6	194	2. 29	1. 45	0. 32
1449	Testis	222205_x_at	Human DNA sequence from clone RP13-329D4 on ch	Hs. 193783. 0	gb:AA446466_RC		UL	2. 6	130	2. 11	1. 26	0. 33
1450	Testis	205747_at	cerebellin 1 precursor	Hs. 662. 0	gb:NM_004352. 1	CBLN1	16	2. 5	143	2. 15	1. 27	0. 36
1451	Testis	206626_x_at	synovial sarcoma, X breakpoint 1	Hs. 194759. 0	gb:BC001003. 2		X	2. 5	179	2. 25	1. 35	0. 36
1452	Testis	208017_s_at	MCF. 2 cell line derived transforming sequence	Hs. 89543. 0	gb:NM_005369. 1	MCF2		2. 5	93	1. 97	0. 94	0. 41
1453	Testis	208541_x_at	transcription factor A, mitochondrial	Hs. 239379. 0	gb:NM_012251. 1	TFAM		2. 5	105	2. 02	1. 23	0. 31
1454	Testis	209743_s_at	atrophin-1 interacting protein 4	Hs. 98074. 0	gb:AI989677_RC		20	2. 5	162	2. 21	1. 64	0. 23
1455	Testis	213038_at	Human DNA sequence from clone RP5-1174N9 on ch	Hs. 64239. 0	gb:AL031602		1	2. 5	367	2. 57	1. 65	0. 36
1456	Testis	219242_at	hypothetical protein FLJ13386	Hs. 300876. 0	gb:NM_025180. 1	FLJ13386	3	2. 5	192	2. 28	1. 40	0. 35
1457	Testis	220270_at	hypothetical protein	Hs. 97464. 0	gb:NM_019038. 1	FLJ11045	13	2. 5	133	2. 12	1. 15	0. 40
1458	Testis	220614_s_at	hypothetical protein FLJ23121	Hs. 98912. 0	gb:NM_024694. 1	FLJ23121	6	2. 5	60	1. 78	0. 65	0. 45
1459	Testis	205449_at	protein predicted by clone 23627	Hs. 23642. 0	gb:NM_013299. 1	HSU79266	11	2. 4	198	2. 30	1. 06	0. 52
1460	Ovary	219331_s_at	hypothetical protein FLJ10748	Hs. 10414. 0	gb:NM_018203. 1	FLJ10748	1	7	644	2. 81	1. 44	0. 19
1461	Ovary	205432_at	oviductal glycoprotein 1, 120kD (mucin 9, ovid	Hs. 1154. 0	gb:NM_002557. 1	OVGP1	1	5. 1	482	2. 68	1. 88	0. 16
1462	Ovary	204581_at	CD22 antigen	Hs. 171763. 0	gb:NM_001771. 1	CD22	19	5	94	1. 97	0. 51	0. 29
1463	Ovary	220102_at	forkhead box L2	Hs. 289292. 0	gb:NM_023067. 1	BPES	3	4. 7	543	2. 73	1. 22	0. 32
1464	Ovary	211062_s_at			gb:BC006393. 1		4	4. 4	291	2. 46	1. 15	0. 30
1465	Ovary	215506_s_at	ras homolog gene family, member I	Hs. 194695. 1	gb:AK021882. 1		1	3. 6	483	2. 68	1. 44	0. 34
1466	Ovary	220911_s_at	KIAA1305 protein	Hs. 288348. 0	gb:NM_025081. 1	KIAA1305	14	3	267	2. 43	1. 56	0. 29
1467	Placenta	208069_x_at	growth hormone 1	Hs. 115352. 2	gb:NM_022561. 1	GH1	17	16. 3	3713	3. 57	0. 84	0. 17
1468	Placenta	211151_x_at	growth hormone 1	Hs. 115352. 5	gb:AF185611. 1	GHV	17	15. 8	3227	3. 51	0. 94	0. 16
1469	Placenta	208356_x_at	chorionic somatomammotropin hormone 1 (placenta	Hs. 324928. 1	gb:NM_022642. 1	CSH1	17	13. 4	7916	3. 90	0. 61	0. 24
1470	Placenta	206885_x_at	growth hormone 1	Hs. 115352. 0	gb:NM_022559. 1	GH1	17	12	4359	3. 64	0. 92	0. 23
1471	Placenta	203807_x_at	chorionic somatomammotropin hormone 2	Hs. 306155. 3	gb:NM_020991. 2	CSH2	17	11. 7	6549	3. 82	0. 43	0. 29
1472	Placenta	208068_x_at	growth hormone 1	Hs. 115352. 3	gb:NM_022562. 1	GH1		11. 7	3940	3. 60	0. 95	0. 23
1473	Placenta	208257_x_at	pregnancy specific beta-1-glycoprotein 1	Hs. 173609. 0	gb:NM_006905. 1	PSG1	19	11. 5	1179	3. 07	0. 93	0. 19
1474	Placenta	206859_s_at	progestagen-associated endometrial protein (pla	Hs. 82269. 0	gb:NM_002571. 1	PAEP	9	10. 8	263	2. 42	0. 83	0. 15
1475	Placenta	206475_x_at	chorionic somatomammotropin hormone 1 (placenta	Hs. 324928. 3	gb:NM_022640. 1	CSH1	17	10. 5	5699	3. 76	0. 83	0. 28
1476	Placenta	208341_x_at	chorionic somatomammotropin hormone 2	Hs. 306155. 0	gb:NM_022646. 1	CSH2	17	10. 5	6639	3. 82	0. 89	0. 28
1477	Placenta	209594_x_at	pregnancy specific beta-1-glycoprotein 9	Hs. 272620. 1	gb:M34421. 1	PSG1	19	10. 5	955	2. 98	2. 01	0. 09
1478	Placenta	211739_x_at			gb:BC005921. 1		17	10. 2	7057	3. 85	0. 47	0. 33
1479	Placenta	204664_at	alkaline phosphatase, placental (Regan isozyme)	Hs. 284255. 0	gb:NM_001632. 2	ALPP	2	10	146	2. 16	0. 54	0. 16
1480	Placenta	205445_at	prolactin	Hs. 1905. 0	gb:NM_000948. 1	PRL	6	9. 9	1121	3. 05	1. 72	0. 13
1481	Placenta	208342_x_at	chorionic somatomammotropin hormone 2	Hs. 306155. 1	gb:NM_022645. 1	CSH2	17	9. 6	5664	3. 75	0. 81	0. 31
1482	Placenta	215821_x_at	pregnancy specific beta-1-glycoprotein 3	Hs. 282847. 2	gb:R32065		19	9. 6	893	2. 95	1. 95	0. 10

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	A	B	C	D	E	F	G	H	I	J	K	L
1483	Placenta	205958_x_at	chorionic somatomammotropin hormone-like 1	Hs. 287269. 1	gb:NM_022579. 1	CSHL1	17	9. 5	5953	3. 77	0. 56	0. 34
1484	Placenta	208357_x_at	chorionic somatomammotropin hormone 1 (placenta	Hs. 324928. 2	gb:NM_022641. 1	CSH1	17	9. 4	5777	3. 76	0. 60	0. 34
1485	Placenta	202493_x_at	chorionic somatomammotropin hormone 1 (placenta	Hs. 324928. 0	gb:NM_001317. 2	CSH1	17	9. 1	5748	3. 76	0. 50	0. 36
1486	Placenta	205840_x_at	growth hormone 1	Hs. 115352. 4	gb:NM_000515. 2	GH1	17	8. 5	3692	3. 57	1. 13	0. 29
1487	Placenta	206886_x_at	growth hormone 1	Hs. 115352. 1	gb:NM_022560. 1	GH1	17	8. 5	3941	3. 60	0. 87	0. 32
1488	Placenta	201981_at	pregnancy-associated plasma protein A	Hs. 75874. 0	gb:AA148534_RC		9	8. 2	419	2. 62	1. 32	0. 16
1489	Placenta	206519_x_at	sialic acid binding Ig-like lectin 6	Hs. 117992. 0	gb:D86358. 1		19	7. 9	194	2. 29	0. 28	0. 25
1490	Placenta	209738_x_at	pregnancy specific beta-1-glycoprotein 6	Hs. 252097. 1	gb:M31125. 1		19	7. 9	858	2. 93	1. 76	0. 15
1491	Placenta	208294_x_at	chorionic somatomammotropin hormone-like 1	Hs. 287269. 3	gb:NM_022578. 1	CSHL1	17	7. 8	5436	3. 74	0. 82	0. 37
1492	Placenta	207770_x_at	chorionic somatomammotropin hormone 2	Hs. 306155. 2	gb:NM_022644. 1	CSH2	17	7. 3	3753	3. 57	1. 55	0. 28
1493	Placenta	206195_x_at	growth hormone 2	Hs. 65149. 0	gb:NM_022557. 1	GH2	17	7. 2	626	2. 80	0. 90	0. 26
1494	Placenta	208295_x_at	chorionic somatomammotropin hormone-like 1	Hs. 287269. 4	gb:NM_022580. 1	CSHL1	17	7	5540	3. 74	0. 77	0. 42
1495	Placenta	219038_at	hypothetical protein FLJ11565	Hs. 61763. 0	gb:NM_024657. 1	FLJ11565	X	6. 7	627	2. 80	1. 78	0. 15
1496	Placenta	204637_at	glycoprotein hormones, alpha polypeptide	Hs. 119689. 0	gb:NM_000735. 2	CGA	6	6. 6	2246	3. 35	1. 22	0. 32
1497	Placenta	205563_at	KiSS-1 metastasis-suppressor	Hs. 95008. 0	gb:NM_002256. 1	KISS1		6. 5	260	2. 41	0. 68	0. 27
1498	Placenta	207285_x_at	chorionic somatomammotropin hormone-like 1	Hs. 287269. 0	gb:NM_001318. 2	CSHL1	17	6. 5	6540	3. 82	0. 87	0. 45
1499	Placenta	221200_at	PP3227 protein	Hs. 302027. 0	gb:NM_022155. 1	PP3227	14	6. 3	150	2. 18	0. 44	0. 28
1500	Placenta	204702_s_at	nuclear factor (erythroid-derived 2)-like 3	Hs. 22900. 0	gb:NM_004289. 3	NFE2L3	7	6. 1	335	2. 53	1. 44	0. 18
1501	Placenta	204602_at	dickkopf (Xenopus laevis) homolog 1	Hs. 40499. 0	gb:NM_012242. 1	DKK1	10	6	590	2. 77	1. 65	0. 19
1502	Placenta	210287_s_at	fms-related tyrosine kinase 1 (vascular endothe	Hs. 138671. 1	gb:U01134. 1	sflt	13	6	255	2. 41	0. 55	0. 31
1503	Placenta	208134_x_at			gb:NM_031246. 1	PSG2	19	5. 9	1174	3. 07	1. 33	0. 30
1504	Placenta	208293_x_at	chorionic somatomammotropin hormone-like 1	Hs. 287269. 2	gb:NM_022581. 1	CSHL1	17	5. 9	4931	3. 69	0. 89	0. 48
1505	Placenta	205629_s_at	corticotropin releasing hormone	Hs. 75294. 0	gb:BC002599. 1		8	5. 8	114	2. 06	0. 54	0. 26
1506	Placenta	205564_at	JM27 protein	Hs. 95420. 0	gb:NM_007003. 1	JM27	X	5. 7	3571	3. 55	1. 14	0. 42
1507	Placenta	205602_x_at	pregnancy specific beta-1-glycoprotein 7	Hs. 225932. 0	gb:NM_002783. 1	PSG7	19	5. 7	912	2. 96	1. 39	0. 28
1508	Placenta	215729_s_at	TONDU	Hs. 9030. 2	gb:BE542323		X	5. 7	662	2. 82	0. 27	0. 45
1509	Placenta	202952_s_at	a disintegrin and metalloproteinase domain 12 (	Hs. 8850. 0	gb:NM_003474. 2	ADAM12	10	5. 5	945	2. 98	0. 92	0. 37
1510	Placenta	203768_s_at	steroid sulfatase (microsomal), arylsulfatase C	Hs. 79876. 0	gb:AU138166		X	5. 5	197	2. 30	1. 59	0. 13
1511	Placenta	205487_s_at	TONDU	Hs. 9030. 0	gb:NM_016267. 1	TONDU	X	5. 5	278	2. 44	0. 88	0. 29
1512	Placenta	211741_x_at			gb:BC005924. 1		19	5. 5	594	2. 77	1. 38	0. 25
1513	Placenta	204830_x_at	pregnancy specific beta-1-glycoprotein 5	Hs. 251850. 0	gb:NM_002781. 1	PSG5	19	5. 4	684	2. 83	1. 31	0. 28
1514	Placenta	220158_at	placental protein 13-like protein	Hs. 24236. 0	gb:NM_020129. 1	LOC56891	19	5. 4	295	2. 47	1. 81	0. 12
1515	Placenta	219424_at	Epstein-Barr virus induced gene 3	Hs. 185705. 0	gb:NM_005755. 1	EBI3	19	5. 3	465	2. 67	0. 73	0. 37
1516	Placenta	203475_at	cytochrome P450, subfamily XIX (aromatization c	Hs. 79946. 0	gb:NM_000103. 1	CYP19	15	5. 2	985	2. 99	1. 52	0. 28
1517	Placenta	204943_at	a disintegrin and metalloproteinase domain 12 (	Hs. 8850. 1	gb:NM_021641. 1	ADAM12	10	5. 2	1350	3. 13	1. 83	0. 25
1518	Placenta	213894_at	KIAA0960 protein	Hs. 29900. 0	gb:BF447246_RC		7	5. 2	295	2. 47	1. 59	0. 17
1519	Placenta	210367_s_at	prostaglandin E synthase	Hs. 146688. 1	gb:AF010316. 1	PIG12	9	5. 1	319	2. 50	1. 70	0. 16
1520	Placenta	202980_s_at	seven in absentia (Drosophila) homolog 1	Hs. 295923. 0	gb:AI953523_RC		16	5	98	1. 99	1. 40	0. 12

Fig. 81

	A	B	C	D	E	F	G	H	I	J	K	L
1521	Placenta	203770_s_at	steroid sulfatase (microsomal), arylsulfatase (	Hs. 79876. 0	gb:J04964. 1	STS	X	5	71	1. 85	0. 68	0. 23
1522	Placenta	205777_at	dual specificity phosphatase 9	Hs. 144879. 0	gb:NM_001395. 1	DUSP9		5	370	2. 57	1. 55	0. 20
1523	Placenta	203399_x_at	pregnancy specific beta-1-glycoprotein 3	Hs. 282847. 0	gb:NM_021016. 1	PSG3	19	4. 8	574	2. 76	1. 19	0. 33
1524	Placenta	206269_at	glial cells missing (Drosophila) homolog a	Hs. 28346. 0	gb:AB026493. 1			4. 6	268	2. 43	1. 16	0. 28
1525	Placenta	205191_at	retinitis pigmentosa 2 (X-linked recessive)	Hs. 44766. 0	gb:NM_006915. 1	RP2	X	4. 5	122	2. 09	1. 24	0. 19
1526	Placenta	209277_at	tissue factor pathway inhibitor 2	Hs. 295944. 0	gb:AL574096_RC		7	4. 4	1129	3. 05	1. 09	0. 45
1527	Placenta	219955_at	hypothetical protein	Hs. 150890. 0	gb:NM_019079. 1	FLJ10884	1	4. 4	299	2. 48	1. 53	0. 22
1528	Placenta	220267_at	hypothetical protein	Hs. 87383. 0	gb:NM_019016. 1	FLJ20261	17	4. 4	237	2. 37	1. 42	0. 22
1529	Placenta	201982_s_at	pregnancy-associated plasma protein A	Hs. 75874. 0	gb:NM_002581. 1	PAPPA	9	4. 3	284	2. 45	1. 19	0. 30
1530	Placenta	202410_x_at	insulin-like growth factor 2 (somatomedin A)	Hs. 251664. 0	gb:NM_000612. 2	IGF2	NA	4. 3	491	2. 69	0. 96	0. 40
1531	Placenta	207085_x_at	colony stimulating factor 2 receptor, alpha, 1(	Hs. 182378. 0	gb:NM_006140. 1	CSF2RA		4. 3	187	2. 27	0. 82	0. 33
1532	Placenta	211508_s_at	growth hormone 2	Hs. 65149. 1	gb:AF006060. 1	hGH-V	17	4. 3	227	2. 36	1. 16	0. 28
1533	Placenta	219824_at	sulfate transporter 1	Hs. 9098. 0	gb:NM_012450. 1	SUT1		4. 2	319	2. 50	1. 66	0. 20
1534	Placenta	210195_s_at	pregnancy specific beta-1-glycoprotein 1	Hs. 173609. 2	gb:M34715. 1		19	4. 1	217	2. 34	1. 57	0. 19
1535	Placenta	204515_at	hydroxy-delta-5-steroid dehydrogenase, 3 beta-	Hs. 38586. 0	gb:NM_000862. 1	HSD3B1	1	4	1010	3. 00	1. 07	0. 48
1536	Placenta	214702_at	Homo sapiens mRNA for MSF-FN70 (FN gene)	Hs. 321592. 0	gb:AJ276395. 1	FN	2	4	539	2. 73	1. 01	0. 43
1537	Placenta	210889_s_at	Fc fragment of IgG, low affinity IIb, receptor	Hs. 278443. 3	gb:M31933. 1		1	3. 9	857	2. 93	1. 68	0. 32
1538	Placenta	211042_x_at			gb:BC006329. 1			3. 9	444	2. 65	1. 82	0. 21
1539	Placenta	214801_at	Homo sapiens cDNA FLJ11392 fis, clone HEMBA100(	Hs. 288924. 0	gb:W88821		1	3. 8	310	2. 49	1. 94	0. 14
1540	Placenta	205829_at	hydroxysteroid (17-beta) dehydrogenase 1	Hs. 85279. 0	gb:NM_000413. 1	HSD17B1	17	3. 7	591	2. 77	0. 90	0. 51
1541	Placenta	211607_x_at			gb:U48722. 1		7	3. 7	256	2. 41	1. 29	0. 31
1542	Placenta	206549_at	insulin-like 4 (placenta)	Hs. 21666. 0	gb:NM_002195. 1	INSL4	9	3. 6	239	2. 38	1. 59	0. 22
1543	Placenta	203625_x_at	S-phase kinase-associated protein 2 (p45)	Hs. 23348. 0	gb:BG105365		5	3. 5	692	2. 84	1. 63	0. 34
1544	Placenta	204569_at	MAK-related kinase	Hs. 108850. 0	gb:NM_014920. 1	KIAA0936	6	3. 5	89	1. 95	1. 23	0. 21
1545	Placenta	205431_s_at	bone morphogenetic protein 5	Hs. 1104. 0	gb:NM_021073. 1	BMP5	6	3. 4	198	2. 30	1. 75	0. 16
1546	Placenta	219702_at	placenta-specific 1	Hs. 13026. 0	gb:NM_021796. 1	PLAC1	X	3. 4	165	2. 22	1. 58	0. 19
1547	Placenta	209757_s_at	v-myc avian myelocytomatosis viral related onc(	Hs. 25960. 1	gb:BC002712. 1		2	3. 3	116	2. 06	0. 82	0. 37
1548	Placenta	203819_s_at	IGF-II mRNA-binding protein 3	Hs. 79440. 0	gb:AU160004_RC		7	3. 2	178	2. 25	0. 87	0. 44
1549	Placenta	209652_s_at	placental growth factor, vascular endothelial g	Hs. 2894. 1	gb:BC001422. 1		14	3. 2	286	2. 46	1. 04	0. 45
1550	Placenta	210196_s_at	pregnancy specific beta-1-glycoprotein 1	Hs. 173609. 2	gb:M33663. 1		19	3. 2	46	1. 66	0. 46	0. 38
1551	Placenta	220736_at	solute carrier family 19, member 3	Hs. 306236. 0	gb:NM_025243. 1	SLC19A3	2	3. 2	140	2. 15	1. 41	0. 23
1552	Placenta	213790_at	a disintegrin and metalloproteinase domain 12 (	Hs. 8850. 2	gb:W46291_RC		10	3. 1	167	2. 22	1. 13	0. 35
1553	Placenta	215717_s_at	fibrillin 2 (congenital contractural arachnoda(	Hs. 79432. 1	gb:X62009. 1		5	3. 1	136	2. 13	1. 20	0. 30
1554	Placenta	219761_at	C-type lectin-like receptor-1	Hs. 29549. 0	gb:NM_016511. 1	LOC51267	12	3. 1	331	2. 52	1. 81	0. 23
1555	Placenta	213183_s_at	cyclin-dependent kinase inhibitor 1C (p57, Kip2	Hs. 106070. 2	gb:N95363_RC		11	3	228	2. 36	1. 49	0. 28
1556	Placenta	205555_s_at	msh (Drosophila) homeo box homolog 2	Hs. 89404. 0	gb:D31771. 1	HDPMSX2A	5	2. 9	168	2. 22	0. 84	0. 48
1557	Placenta	210274_at	melanoma antigen, family A, 8	Hs. 37109. 0	gb:BC002455. 1		X	2. 9	213	2. 33	1. 17	0. 40
1558	Placenta	214858_at	Homo sapiens clone 24566 mRNA sequence	Hs. 133342. 0	gb:AF070536. 1		2	2. 6	75	1. 88	0. 57	0. 49

Fig. 82

	A	B	C	D	E	F	G	H	I	J	K	L
1559	Placenta	219523_s_at	hypothetical protein FLJ10474	Hs. 41793. 0	gb:NM_018104. 1	FLJ10474	4	2. 5	146	2. 16	1. 33	0. 34
1560	Placenta	219552_at	likely ortholog of mouse polydom	Hs. 271350. 0	gb:NM_024500. 1	POLYDOM	9	2. 5	66	1. 82	0. 85	0. 39
1561	Placenta	202966_at	calpain 6	Hs. 169172. 0	gb:NM_014289. 2	CAPN6	X	2. 4	105	2. 02	0. 88	0. 48
1562	Uterus	205680_at	matrix metalloproteinase 10 (stromelysin 2)	Hs. 2258. 0	gb:NM_002425. 1	MMP10	11	5. 2	953	2. 98	0. 91	0. 40
1563	Uterus	214373_at	protein phosphatase 4 regulatory subunit 2	Hs. 125682. 1	gb:AI582773_RC		3	4. 5	233	2. 37	1. 74	0. 14
1564	Uterus	219993_at	hypothetical protein FLJ22252 similar to SRY-bo	Hs. 97984. 0	gb:NM_022454. 1	FLJ22252	8	4. 2	203	2. 31	1. 71	0. 14
1565	Uterus	205534_at	BH-protocadherin (brain-heart)	Hs. 34073. 0	gb:NM_002589. 1	PCDH7	4	3. 9	441	2. 64	1. 14	0. 39
1566	Uterus	211029_x_at			gb:BC006245. 1			3. 9	53	1. 73	0. 72	0. 26
1567	Uterus	219249_s_at	hypothetical protein FLJ22041 similar to FK506	Hs. 3849. 0	gb:NM_021939. 1	FLJ22041	17	3. 9	150	2. 18	0. 99	0. 30
1568	Uterus	205932_s_at	msh (Drosophila) homeo box homolog 1 (formerly	Hs. 1494. 0	gb:NM_002448. 1	MSX1	4	3. 5	502	2. 70	1. 70	0. 29
1569	Uterus	212344_at	KIAA1077 protein	Hs. 70823. 0	gb:AW043713_RC		8	3. 4	469	2. 67	1. 72	0. 28
1570	Uterus	220399_at	hypothetical protein FLJ22639	Hs. 157184. 0	gb:NM_024796. 1	FLJ22639	6	3. 4	104	2. 02	1. 23	0. 23
1571	Uterus	213707_s_at	distal-less homeo box 5	Hs. 99348. 0	gb:NM_005221. 3	DLX5	7	3. 2	197	2. 29	1. 42	0. 28
1572	Uterus	205535_s_at	BH-protocadherin (brain-heart)	Hs. 34073. 0	gb:NM_002589. 1	PCDH7	4	3	70	1. 84	0. 67	0. 39
1573	Bone marrow	207384_at	peptidoglycan recognition protein	Hs. 137583. 0	gb:NM_005091. 1	PGLYRP	19	17	963	2. 98	0. 56	0. 14
1574	Bone marrow	207269_at	defensin, alpha 4, corticostatin	Hs. 2582. 0	gb:NM_001925. 1	DEFA4	8	15. 1	7263	3. 86	1. 66	0. 15
1575	Bone marrow	203949_at	myeloperoxidase	Hs. 1817. 0	gb:NM_000250. 1	MPO	17	13. 7	2609	3. 42	1. 55	0. 14
1576	Bone marrow	206111_at	ribonuclease, RNase A family, 2 (liver, eosinop	Hs. 728. 0	gb:NM_002934. 1	RNASE2	14	10. 7	3951	3. 60	1. 62	0. 18
1577	Bone marrow	210504_at	Kruppel-like factor 1 (erythroid)	Hs. 37860. 0	gb:U65404. 1		19	10. 3	513	2. 71	1. 43	0. 12
1578	Bone marrow	210746_s_at	erythrocyte membrane protein band 4. 2	Hs. 733. 1	gb:M30646. 1		15	9. 8	1456	3. 16	0. 98	0. 22
1579	Bone marrow	205592_at	solute carrier family 4, anion exchanger, membe	Hs. 185923. 0	gb:X77737. 1		17	9. 2	3726	3. 57	1. 13	0. 26
1580	Bone marrow	206937_at	spectrin, alpha, erythrocytic 1 (elliptocytosis	Hs. 1985. 0	gb:NM_003126. 1	SPTA1	1	9. 2	441	2. 64	1. 39	0. 14
1581	Bone marrow	214575_s_at	azurocidin 1 (cationic antimicrobial protein 37	Hs. 72885. 0	gb:NM_001700. 1	AZU1	NA	8. 9	5646	3. 75	1. 17	0. 29
1582	Bone marrow	203948_s_at	myeloperoxidase	Hs. 1817. 0	gb:J02694. 1	MPO	17	8. 8	3891	3. 59	0. 62	0. 34
1583	Bone marrow	206207_at	Charot-Leyden crystal protein	Hs. 889. 0	gb:NM_001828. 3	CLC	19	8. 8	3877	3. 59	1. 56	0. 23
1584	Bone marrow	221627_at	ring finger protein 9	Hs. 274295. 1	gb:AF220123. 1	TRIM10	6	8. 7	654	2. 82	1. 89	0. 11
1585	Bone marrow	211560_s_at	cytochrome b5 outer mitochondrial membrane prec	Hs. 79103. 1	gb:AF130113. 1			8. 6	1888	3. 28	0. 82	0. 29
1586	Bone marrow	219672_at	erythroid differentiation-related factor	Hs. 274309. 0	gb:NM_016633. 1	LOC51327		8. 6	2125	3. 33	1. 65	0. 20
1587	Bone marrow	206871_at	elastase 2, neutrophil	Hs. 99863. 0	gb:NM_001972. 1	ELA2	2	8. 4	6800	3. 83	0. 90	0. 35
1588	Bone marrow	206283_s_at	T-cell acute lymphocytic leukemia 1	Hs. 73828. 0	gb:NM_003189. 1	TAL1	1	8. 3	587	2. 77	1. 70	0. 13
1589	Bone marrow	207341_at	proteinase 3 (serine proteinase, neutrophil, We	Hs. 928. 0	gb:NM_002777. 2	PRTN3	NA	7. 2	2333	3. 37	1. 38	0. 28
1590	Bone marrow	220811_at	proteoglycan 3	Hs. 251386. 0	gb:NM_006093. 2	PRG3	11	7. 2	2114	3. 33	1. 52	0. 25
1591	Bone marrow	216833_x_at	Human glycophorin HeP2 mRNA, partial cds	Hs. 307185. 0	gb:U05255. 1			6. 9	616	2. 79	1. 50	0. 19
1592	Bone marrow	211821_x_at	glycophorin A (includes MN blood group)	Hs. 108694. 1	gb:U00178. 1	GPErik		6. 8	539	2. 73	0. 60	0. 31
1593	Bone marrow	206851_at	ribonuclease, RNase A family, 3 (eosinophil cat	Hs. 73839. 0	gb:NM_002935. 1	RNASE3	14	6. 7	3386	3. 53	1. 32	0. 33
1594	Bone marrow	205653_at	cathepsin G	Hs. 100764. 0	gb:NM_001911. 1	CTSG	14	6. 6	3003	3. 48	1. 49	0. 30
1595	Bone marrow	204788_s_at	protoporphyrinogen oxidase	Hs. 100016. 0	gb:NM_000309. 1	PPOX		6. 3	293	2. 47	1. 90	0. 09
1596	Bone marrow	206390_x_at	platelet factor 4	Hs. 81564. 0	gb:NM_002619. 1	PF4	4	6	1302	3. 11	1. 77	0. 22

Fig. 83

	A	B	C	D	E	F	G	H	I	J	K	L
1597	Bone marrow	216925_s_at	T-cell acute lymphocytic leukemia 1	Hs. 73828. 1	gb:X51990. 1		1	5. 8	258	2. 41	1. 29	0. 20
1598	Bone marrow	221764_at	glycerol-3-phosphate dehydrogenase 1 (soluble)	Hs. 9739. 2	gb:AL574186_RC		2	5. 8	791	2. 90	2. 00	0. 15
1599	Bone marrow	211254_x_at	Rhesus blood group-associated glycoprotein	Hs. 169536. 2	gb:AF031549. 1	Rh50	6	5. 6	450	2. 65	2. 04	0. 11
1600	Bone marrow	206676_at	carcinoembryonic antigen-related cell adhesion	Hs. 41. 0	gb:M33326. 1	NCA	19	5. 5	2218	3. 35	0. 64	0. 49
1601	Bone marrow	220001_at	peptidyl arginine deiminase, type V	Hs. 117232. 0	gb:NM_012387. 1	PAD	1	5. 4	842	2. 93	0. 98	0. 36
1602	Bone marrow	203115_at	ferrochelatase (protoporphyria)	Hs. 26. 0	gb:AU152635_RC		18	5. 3	243	2. 39	1. 50	0. 17
1603	Bone marrow	218978_s_at	hypothetical protein PRO1584	Hs. 283716. 0	gb:NM_018586. 1	PRO1584	8	5. 3	202	2. 30	1. 49	0. 15
1604	Bone marrow	214627_at	eosinophil peroxidase	Hs. 46295. 0	gb:X14346. 1		17	5. 2	1159	3. 06	1. 78	0. 24
1605	Bone marrow	207459_x_at	glycophorin B (includes Ss blood group)	Hs. 250653. 0	gb:NM_002100. 2	GYPB	4	5	782	2. 89	1. 62	0. 26
1606	Bone marrow	210254_at	membrane-spanning 4-domains, subfamily A, membe	Hs. 99960. 0	gb:L35848. 1	HTm4	11	5	1242	3. 09	0. 83	0. 45
1607	Bone marrow	211413_s_at	Homo sapiens PADI-H protein mRNA, complete cds	Hs. 283960. 0	gb:AF229067. 1		1	5	199	2. 30	0. 40	0. 38
1608	Bone marrow	215037_s_at	BCL2-like 1	Hs. 305890. 2	gb:U72398		20	5	231	2. 36	1. 63	0. 15
1609	Bone marrow	206660_at	immunoglobulin lambda-like polypeptide 1	Hs. 288168. 0	gb:NM_020070. 1	IGLL1	22	4. 9	314	2. 50	1. 23	0. 26
1610	Bone marrow	207765_s_at	hypothetical protein FLJ11560	Hs. 301696. 0	gb:NM_025182. 1	FLJ11560	9	4. 7	240	2. 38	1. 77	0. 13
1611	Bone marrow	211820_x_at	glycophorin A (includes MN blood group)	Hs. 108694. 2	gb:U00179. 1	GPErik		4. 7	411	2. 61	0. 78	0. 39
1612	Bone marrow	206145_at	Rhesus blood group-associated glycoprotein	Hs. 169536. 0	gb:NM_000324. 1	RHAG	6	4. 6	474	2. 68	1. 00	0. 36
1613	Bone marrow	210446_at	GATA-binding protein 1 (globin transcription fa	Hs. 765. 0	gb:M30601. 1		X	4. 6	180	2. 26	1. 41	0. 18
1614	Bone marrow	56748_at						4. 5	89	1. 95	1. 32	0. 14
1615	Bone marrow	204768_s_at	flap structure-specific endonuclease 1	Hs. 4756. 0	gb:NM_004111. 3	FEN1	11	4. 5	224	2. 35	1. 76	0. 13
1616	Bone marrow	205936_s_at	hexokinase 3 (white cell)	Hs. 159237. 0	gb:NM_002115. 1	HK3	5	4. 5	426	2. 63	0. 91	0. 38
1617	Bone marrow	211307_s_at	Fc fragment of IgA, receptor for	Hs. 193122. 3	gb:U43677. 1	CD89	19	4. 5	111	2. 05	0. 40	0. 37
1618	Bone marrow	214407_x_at	glycophorin B (includes Ss blood group)	Hs. 250653. 1	gb:AI240545_RC		4	4. 5	940	2. 97	1. 45	0. 34
1619	Bone marrow	204859_s_at	apoptotic protease activating factor	Hs. 77579. 1	gb:NM_013229. 1	APAF1	12	4. 4	93	1. 97	1. 48	0. 11
1620	Bone marrow	206371_at	folate receptor 3 (gamma)	Hs. 352. 0	gb:NM_000804. 2	FOLR3	11	4. 4	280	2. 45	1. 24	0. 28
1621	Bone marrow	210044_s_at	lymphoblastic leukemia derived sequence 1	Hs. 46446. 0	gb:BC002796. 1		19	4. 3	383	2. 58	1. 13	0. 34
1622	Bone marrow	215047_at	DKFZP434C091 protein	Hs. 51692. 0	gb:AL080170. 1	DKFZp434C0	1	4. 3	712	2. 85	0. 40	0. 57
1623	Bone marrow	215819_s_at	Rhesus blood group, CcEe antigens	Hs. 278994. 1	gb:N53959_RC		1	4. 3	542	2. 73	1. 62	0. 26
1624	Bone marrow	216317_x_at	Rhesus blood group, CcEe antigens	Hs. 278994. 2	gb:X63095. 1		1	4. 3	595	2. 77	1. 31	0. 34
1625	Bone marrow	216667_at	Human ECRP gene for eosinophil cationic relate	Hs. 166037. 0	gb:X55989			4. 2	310	2. 49	1. 20	0. 31
1626	Bone marrow	204961_s_at	neutrophil cytosolic factor 1 (47kD, chronic gr	Hs. 1583. 0	gb:NM_000265. 1	NCF1	7	4. 1	541	2. 73	0. 62	0. 51
1627	Bone marrow	209930_s_at	nuclear factor (erythroid-derived 2), 45kD	Hs. 75643. 1	gb:L13974. 1		12	4. 1	494	2. 69	0. 98	0. 42
1628	Bone marrow	205627_at	cytidine deaminase	Hs. 72924. 0	gb:NM_001785. 1	CDA	1	4	672	2. 83	1. 10	0. 44
1629	Bone marrow	210430_x_at	Rhesus blood group, D antigen	Hs. 283822. 0	gb:L08429. 1		1	4	321	2. 51	1. 21	0. 32
1630	Bone marrow	220570_at	found in inflammatory zone 3	Hs. 283091. 0	gb:NM_020415. 2	FIZZ3	19	4	744	2. 87	1. 14	0. 43
1631	Bone marrow	205268_s_at	adducin 2 (beta)	Hs. 247423. 6	gb:NM_017488. 1	ADD2	2	3. 9	310	2. 49	1. 11	0. 36
1632	Bone marrow	205570_at	phosphatidylinositol-4-phosphate 5-kinase, type	Hs. 108966. 0	gb:NM_005028. 2	PIP5K2A	10	3. 8	250	2. 40	1. 51	0. 24
1633	Bone marrow	207854_at	glycophorin E	Hs. 93223. 0	gb:NM_002102. 1	GYPE	4	3. 8	67	1. 83	0. 40	0. 38
1634	Bone marrow	203470_s_at	pleckstrin	Hs. 77436. 0	gb:AI433595_RC		2	3. 6	735	2. 87	1. 49	0. 38

Fig. 84

	A	B	C	D	E	F	G	H	I	J	K	L
1635	Bone marrow	206247_at	MHC class I polypeptide-related sequence B	Hs. 211580. 0	gb:NM_005931. 1	MICB	6	3. 6	200	2. 30	1. 59	0. 20
1636	Bone marrow	206665_s_at	BCL2-like 1	Hs. 305890. 0	gb:NM_001191. 1	BCL2L1	20	3. 6	102	2. 01	0. 69	0. 37
1637	Bone marrow	214523_at	CCAATenhancer binding protein (CEBP), epsilon	Hs. 158323. 0	gb:NM_001805. 1	CEBPE	14	3. 6	164	2. 21	1. 16	0. 29
1638	Bone marrow	215435_at	Homo sapiens cDNA FLJ11921 fis, clone HEMBB100(	Hs. 306616. 0	gb:AK021983. 1_RC		4	3. 6	119	2. 07	1. 46	0. 17
1639	Bone marrow	37145_at						3. 4	371	2. 57	1. 15	0. 41
1640	Bone marrow	205715_at	bone marrow stromal cell antigen 1	Hs. 169998. 0	gb:NM_004334. 1	BST1	4	3. 4	267	2. 43	1. 04	0. 41
1641	Bone marrow	211713_x_at			gb:BC005832. 1		15	3. 4	248	2. 39	1. 68	0. 21
1642	Bone marrow	214316_x_at	calreticulin	Hs. 16488. 3	gb:AI378706			3. 4	79	1. 90	1. 42	0. 14
1643	Bone marrow	204826_at	cyclin F	Hs. 1973. 0	gb:NM_001761. 1	CCNF		3. 2	154	2. 19	1. 15	0. 32
1644	Bone marrow	202856_s_at	solute carrier family 16 (monocarboxylic acid t	Hs. 85838. 0	gb:NM_004207. 1	SLC16A3	17	3. 1	302	2. 48	1. 74	0. 24
1645	Bone marrow	217573_at	ESTs, Highly similar to N-methyl-D-aspartate re	Hs. 299882. 0	gb:AW138468_RC			2. 9	93	1. 97	0. 95	0. 35
1646	Bone marrow	205180_s_at	a disintegrin and metalloproteinase domain 8	Hs. 86947. 0	gb:NM_001109. 1	ADAM8	10	2. 6	177	2. 25	1. 22	0. 40
1647	Bone marrow	210429_at	Rhesus blood group, D antigen	Hs. 283822. 0	gb:X63097. 1		1	2. 4	51	1. 71	0. 90	0. 34
1648	Fetal Brain	205347_s_at	thymosin, beta, identified in neuroblastoma cel	Hs. 56145. 0	gb:NM_021992. 1	TMSNB	X	9. 5	1224	3. 09	1. 76	0. 14
1649	Fetal Brain	204914_s_at	SRY (sex determining region Y)-box 11	Hs. 32964. 0	gb:AW157202_RC		2	6. 2	1575	3. 20	0. 84	0. 38
1650	Fetal Brain	204819_at	faciogenital dysplasia (Aarskog-Scott syndrome)	Hs. 1572. 0	gb:NM_004463. 1	FGD1	X	5. 7	310	2. 49	1. 84	0. 11
1651	Fetal Brain	207658_s_at	forkhead box G1A	Hs. 169277. 0	gb:NM_004471. 1	FOXG1A	14	5. 7	488	2. 69	0. 90	0. 31
1652	Fetal Brain	204851_s_at	doublecortex; lissencephaly, X-linked (doublecd	Hs. 34780. 0	gb:AF040254. 1	DCX	X	5. 4	1345	3. 13	0. 65	0. 46
1653	Fetal Brain	204913_s_at	SRY (sex determining region Y)-box 11	Hs. 32964. 0	gb:AI360875_RC		2	5. 2	1416	3. 15	1. 09	0. 40
1654	Fetal Brain	204915_s_at	SRY (sex determining region Y)-box 11	Hs. 32964. 0	gb:AB028641. 1	sox11	2	5. 1	946	2. 98	0. 90	0. 41
1655	Fetal Brain	211391_s_at	zinc finger protein 278	Hs. 27801. 0	gb:AF254087. 1	EWSZSG fusion		4. 9	218	2. 34	1. 74	0. 12
1656	Fetal Brain	215119_at	KIAA0865 protein	Hs. 175411. 0	gb:AI522028_RC		13	4. 8	409	2. 61	0. 85	0. 36
1657	Fetal Brain	202921_s_at	ankyrin 2, neuronal	Hs. 117970. 0	gb:NM_001148. 2	ANK2	4	4. 5	297	2. 47	1. 86	0. 14
1658	Fetal Brain	214051_at	thymosin, beta, identified in neuroblastoma cel	Hs. 56145. 1	gb:BF677486		X	4. 3	170	2. 23	1. 70	0. 12
1659	Fetal Brain	206103_at	ras-related C3 botulinum toxin substrate 3 (rhd	Hs. 45002. 0	gb:NM_005052. 1	RAC3		4. 2	109	2. 04	0. 85	0. 28
1660	Fetal Brain	207164_s_at	zinc finger protein 238	Hs. 69997. 0	gb:NM_006352. 1	ZNF238	1	4. 2	398	2. 60	1. 02	0. 37
1661	Fetal Brain	220025_at	T-box, brain, 1	Hs. 210862. 0	gb:NM_006593. 1	TBR1	2	4. 1	357	2. 55	1. 54	0. 25
1662	Fetal Brain	204996_s_at	cyclin-dependent kinase 5, regulatory subunit 1	Hs. 93597. 0	gb:NM_003885. 1	CDK5R1	17	4	150	2. 18	0. 92	0. 32
1663	Fetal Brain	206972_s_at	G-protein coupled receptor	Hs. 301642. 0	gb:NM_007369. 1	RE2	1	3. 8	108	2. 03	0. 69	0. 36
1664	Fetal Brain	207147_at	distal-less homeo box 2	Hs. 419. 0	gb:NM_004405. 2	DLX2	2	3. 8	276	2. 44	0. 90	0. 41
1665	Fetal Brain	214577_at	microtubule-associated protein 1B	Hs. 103042. 0	gb:BG164365		5	3. 7	395	2. 60	1. 31	0. 35
1666	Fetal Brain	220277_at	Dvl-binding protein IDAX (inhibition of the Dv	Hs. 118569. 0	gb:NM_025212. 1	IDAX	4	3. 7	219	2. 34	1. 49	0. 23
1667	Fetal Brain	208121_s_at			gb:NM_002848. 2	PTPRO	12	3. 5	462	2. 66	1. 66	0. 28
1668	Fetal Brain	215045_at	Homo sapiens, clone IMAGE:3509783, mRNA, parti	Hs. 332322. 0	gb:BC004145. 1			3. 5	260	2. 41	1. 31	0. 32
1669	Fetal Brain	203228_at	platelet-activating factor acetylhydrolase, isd	Hs. 6793. 0	gb:NM_002573. 1	PAFAH1B3	19	3. 4	547	2. 74	1. 21	0. 45
1670	Fetal Brain	206436_at	chromosome 22 open reading frame 1	Hs. 159538. 0	gb:NM_001585. 1	C22ORF1	22	3. 4	390	2. 59	1. 46	0. 33
1671	Fetal Brain	219668_at	hypothetical protein MGC3129 similar to ganglid	Hs. 20977. 0	gb:NM_024034. 1	MGC3129	20	3. 4	206	2. 31	1. 13	0. 35
1672	Fetal Brain	205536_at	vav 2 oncogene	Hs. 4248. 0	gb:NM_003371. 1	VAV2	9	3. 3	72	1. 85	0. 95	0. 28

Fig. 85

	A	B	C	D	E	F	G	H	I	J	K	L
1673	Fetal Brain	210933_s_at	Homo sapiens, clone MGC:4655, mRNA, complete cd	Hs. 326737. 0	gb:BC004908. 1		16	3. 3	356	2. 55	1. 53	0. 31
1674	Fetal Brain	209757_s_at	v-myc avian myelocytomatosis viral related onc	Hs. 25960. 1	gb:BC002712. 1		2	3. 2	321	2. 51	0. 97	0. 49
1675	Fetal Brain	219742_at	hypothetical protein MGC10772	Hs. 130316. 0	gb:NM_030567. 1	MGC10772	5	3. 2	108	2. 03	0. 91	0. 35
1676	Fetal Brain	206326_at	gastrin-releasing peptide	Hs. 1473. 0	gb:NM_002091. 1	GRP	18	3. 1	331	2. 52	1. 11	0. 46
1677	Fetal Brain	207084_at	POU domain, class 3, transcription factor 2	Hs. 182505. 0	gb:NM_005604. 1	POU3F2	6	3. 1	133	2. 12	0. 90	0. 39
1678	Fetal Brain	219740_at	hypothetical protein FLJ12505	Hs. 125741. 0	gb:NM_024749. 1	FLJ12505	1	2. 5	77	1. 89	0. 95	0. 38
1679	Fetal Liver	206647_at	hemoglobin, zeta	Hs. 272003. 0	gb:NM_005332. 2	HBZ	16	9. 7	4534	3. 66	0. 38	0. 34
1680	Fetal Liver	205838_at	glycophorin A (includes MN blood group)	Hs. 108694. 0	gb:NM_002099. 2	GYPA	4	7. 2	754	2. 88	0. 94	0. 27
1681	Fetal Liver	205837_s_at	glycophorin A (includes MN blood group)	Hs. 108694. 0	gb:BC005319. 1		4	6. 7	715	2. 85	-0. 05	0. 43
1682	Fetal Liver	210677_at	sterol O-acyltransferase 2	Hs. 20580. 0	gb:AF059203. 1		12	5. 7	475	2. 68	1. 07	0. 28
1683	Fetal Liver	205919_at	hemoglobin, epsilon 1	Hs. 117848. 0	gb:NM_005330. 2	HBE1	NA	5. 6	3803	3. 58	1. 31	0. 40
1684	Fetal Liver	206146_s_at	Rhesus blood group-associated glycoprotein	Hs. 169536. 0	gb:AF178841. 1		6	5. 5	65	1. 81	0. 53	0. 23
1685	Fetal Liver	215811_at	Homo sapiens clone GLSH-3 similar to gliadin mR	Hs. 275706. 0	gb:AF238870. 1_RC		4	5. 3	203	2. 31	1. 77	0. 10
1686	Fetal Liver	213092_x_at	KIAA0974 protein	Hs. 44131. 2	gb:AW241779_RC		10	4. 8	117	2. 07	1. 45	0. 13
1687	Fetal Liver	204624_at	ATPase, Cu++ transporting, beta polypeptide (Wi	Hs. 84999. 0	gb:NM_000053. 1	ATP7B	13	4. 4	328	2. 52	1. 88	0. 15
1688	Fetal Liver	219555_s_at	uncharacterized bone marrow protein BM039	Hs. 283532. 0	gb:NM_018455. 1	BM039	16	4	111	2. 05	0. 67	0. 34
1689	Fetal Liver	202422_s_at	fatty-acid-Coenzyme A ligase, long-chain 4	Hs. 81452. 0	gb:NM_022977. 1	FACL4	X	3. 7	372	2. 57	1. 56	0. 27
1690	Fetal Liver	203027_s_at	mevalonate (diphospho) decarboxylase	Hs. 3828. 0	gb:AI189359_RC		4	3. 7	83	1. 92	0. 61	0. 35
1691	Fetal Liver	206077_at	Kell blood group	Hs. 157. 0	gb:NM_000420. 1	KEL	7	3. 6	337	2. 53	1. 35	0. 33
1692	Fetal Liver	218875_s_at	F-box only protein 5	Hs. 272027. 0	gb:NM_012177. 1	FBXO5	6	3. 6	108	2. 03	1. 18	0. 24
1693	Fetal Liver	203764_at	KIAA0008 gene product	Hs. 77695. 0	gb:NM_014750. 1	KIAA0008	14	3. 5	291	2. 46	1. 23	0. 35
1694	Fetal Liver	207345_at	follistatin	Hs. 9914. 0	gb:NM_006350. 2	FST	5	3. 4	167	2. 22	1. 50	0. 21
1695	Fetal Liver	219546_at	hypothetical protein DKFZp434P0116	Hs. 20137. 0	gb:NM_017593. 1	DKFZp434P0	4	3. 4	123	2. 09	0. 96	0. 33
1696	Fetal Liver	212022_s_at	antigen identified by monoclonal antibody Ki-67	Hs. 80976. 1	gb:BF001806_RC		10	3. 3	356	2. 55	1. 21	0. 41
1697	Fetal Liver	217880_at	cell division cycle 27	Hs. 172405. 0	gb:N21397_RC		17	3. 3	225	2. 35	1. 65	0. 22
1698	Fetal Liver	207194_s_at	intercellular adhesion molecule 4, Landsteiner-	Hs. 108287. 1	gb:NM_001544. 2	ICAM4	19	3. 3	169	2. 23	0. 85	0. 41
1699	Fetal Liver	218662_s_at	chromosome condensation protein G	Hs. 193602. 0	gb:NM_022346. 1	HCAP-G	4	3. 3	150	2. 17	1. 27	0. 27
1700	Fetal Liver	205848_at	growth arrest-specific 2	Hs. 129818. 0	gb:NM_005256. 1	GAS2	11	3. 2	359	2. 56	1. 32	0. 39
1701	Fetal Liver	218663_at	chromosome condensation protein G	Hs. 193602. 0	gb:NM_022346. 1	HCAP-G	4	3. 2	142	2. 15	0. 39	0. 56
1702	Fetal Liver	222074_at	uroporphyrinogen decarboxylase	Hs. 78601. 1	gb:AW614435_RC		1	3. 2	109	2. 04	0. 97	0. 34
1703	Fetal Liver	221521_s_at	HSPC037 protein	Hs. 108196. 1	gb:BC003186. 1		16	3. 1	190	2. 28	0. 89	0. 44
1704	Fetal Liver	205194_at	phosphoserine phosphatase	Hs. 56407. 0	gb:NM_004577. 1	PSPH	7	3	97	1. 99	1. 33	0. 22